

Figure 1. Phage-display selects out the OSCP. Semiquantitative PCR using primers specific for the OSCP sequence reveals enrichment of clones encoding the OSCP with each round of selection using immobilized Bz-423.

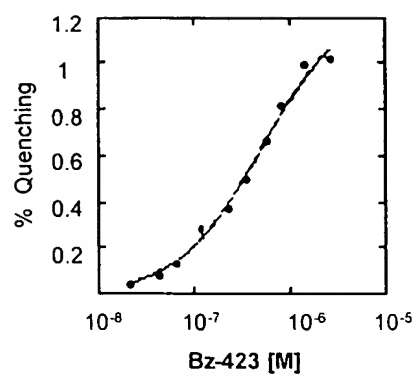


Figure 2. Binding isotherm of Bz-423 and purified human OSCP .

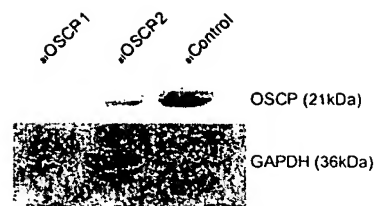


Figure 3. siRNA reduces the OSCP. 10^6 HEK 293 cells transfected with 0.84 μ g of siRNA duplexes specific for OSCP or scrambled control. After 72 h, protein expression determined by immunoblotting whole cell extracts. OSCP is reduced and GAPDH increased as glycolytic response.

5

10

15

FIGURE 4A.

gene	NCBI Accession Number	FOLD INCREASE
glyceraldehyde-3-phosphate dehydrogenase	M33197	23.02
ribosomal protein L29	NM_000892	21.62
heat shock 90kD protein 1, beta	A1216218	21.27
lysosomal-associated multispinning membrane protein-5	NM_006782	20.19
ferritin, light polypeptide	BC538564	19.50
Consensus includes gb:AJ274937.1 /DEF=Homo sapiens partial mRNA for human Ig lambda light chain variable region, clone MB91 (331 bp). /FEA=mRNA /GEN=IGLV /PROI AJ24377		18.95
Consensus includes gb:D84143.1 /DEF=human immunoglobulin (mAb59) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V-J n D84143		18.76
gb:AJ25092.1 /DEF=Homo sapiens mRNA for single-chain antibody, complete cds. /FEA=CDS /PROD=immunoglobulin /DB_XREF=gi:3090425 /UG=Hs.249245 Homo sapiens	AJ25092	18.62
ribosomal protein L29	BF683426	18.53
Consensus includes gb:L48784 /DEF=050 Homo sapiens cDNA /FEA=mRNA /DB_XREF=gi:1068715 /UG=Hs.182426 ribosomal protein S2	L48784	17.74
profilin 1	NM_005022	17.74
ornithine decarboxylase antizyme 1	AF090094	16.18
serine hydroxymethyltransferase 2 (mitochondrial)	NM_005412	15.75
chaperonin containing TCP1, subunit 7 (eta)	NM_006429	15.12
ribosomal protein L6	NM_000973	14.94
macrophage migration inhibitory factor (glycosylation-inhibiting factor)	NM_002415	14.75
Consensus includes gb:D84140.1 /DEF=human immunoglobulin (mAb56) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V-J n D84140		14.48
transketolase (Wernicke-Korsakoff syndrome)	BF696840	14.41
ribosomal protein L18	NM_000979	14.38
transketolase (Wernicke-Korsakoff syndrome)	L12711	13.75
ubiquitin-conjugating enzyme E21 (UBC9 homolog, yeast)	AA910614	12.75
glyceraldehyde-3-phosphate dehydrogenase	M33197	12.69
glyceraldehyde-3-phosphate dehydrogenase	BF689355	12.67
nuclease sensitive element binding protein 1	BE966374	12.57
neutrophil cytosolic factor 4 (40kD)	NM_013416	12.52
T-cell leukemia/lymphoma 1A	X82240	12.46
ribosomal protein L18a	NM_000980	12.33
glyceraldehyde-3-phosphate dehydrogenase	BE561479	12.29
gb:L07950.1 /DEF=Homo sapiens MHC class I HLA B71 mRNA, complete cds. /FEA=CDS /GEN=HLA-B /PROD=MHC HLA B71 /DB_XREF=gi:307236 /FL=gb:L07950.1	L07950	12.26
ubiquitin carrier protein	NM_014501	12.12
actin, beta	X00351	11.85
enolase 1, (alpha)	U88668	11.53
replication factor C (activator 1) 2 (40kD)	M87338	11.48
phosphoserine aminotransferase	NM_021154	10.95
MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)	AF279900	10.74
ribosomal protein S5	NM_001009	10.69
RNA polymerase II transcriptional regulation mediator (Med6, S. cerevisiae, homolog of)	NM_005466	10.68
eukaryotic translation initiation factor 3, subunit 8 (110kD)	BC000533	10.65
chaperonin containing TCP1, subunit 2 (beta)	AL545982	10.52
CGI-135 protein	NM_016068	10.35
polymerase (DNA directed), delta 2, regulatory subunit (50kD)	NM_006230	10.25
ribosomal protein L13	AW574664	9.93
procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone binding protein p55)	J02763	9.85
HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae)	NM_001536	9.83
cold inducible RNA binding protein	NM_001280	9.66
ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit	BE798517	9.57
pinin, desmosome associated protein	AF112222	9.50
pyruvate kinase, muscle	NM_002654	9.43
gb:L23516.1 /DEF=Human Ig rearranged gamma-chain, V-DXP4-JH6c, complete cds. /FEA=mRNA /DB_XREF=gi:385218 /FL=gb:L23516.1	L23516	9.42
fusion, derived from t(12;16) malignant liposarcoma	NM_004960	9.37
immunoglobulin heavy constant mu	U80139	9.32
maternal G10 transcript	NM_003910	9.30
major histocompatibility complex, class I, B	D83043	9.29
small EDRK-kch factor 2	NM_005770	9.29
heat shock 90kD protein 1, beta	AF275719	9.05
Consensus includes gb:AF254822 /DEF=Homo sapiens SMARCA4 isoform (SMARCA4) gene, complete cds, alternatively spliced /FEA=CDS_2 /DB_XREF=gi:10946127 /UG=		8.88
protein phosphatase 1G (formerly 2C), magnesium-dependent, gamma isoform	NM_002707	8.77
glucose phosphate isomerase	NM_000175	8.76
proteasome (prosome, macropain) 26S subunit, non-ATPase, 8	NM_002812	8.74

ribosomal protein S19	BE259729	8.68
eukaryotic translation elongation factor 1 alpha 1	AL035687	8.48
gbL23518.1 /DEF=Human Ig rearranged gamma-chain, V-DXP1-JH4b, complete cds. /FEA=mRNA /DB_XREF=g:385220 /FL=gb:L23518.1	L23518	8.41
ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit	NM_001687	8.41
procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone binding protein p55)	NM_000918	8.38
endothelial differentiation-related factor 1	AB002282	8.37
acid phosphatase 1, soluble	BG035989	8.33
chaperonin containing TCP1, subunit 3 (gamma)	NM_005988	8.32
proteasome (prosome, macropain) subunit, alpha type, 7	AL078633	8.23
Consensus includes gb:AF043584.1 /DEF=Homo sapiens clone ASMneg1-b1 Immunoglobulin lambda chain VJ region, (IGL) mRNA, partial cds. /FEA=mRNA /GEN=IGL /PRO AF043584	AF043584	8.21
non-POU-domain-containing, octamer-binding	L14599	8.09
CGI-51 protein	NM_015380	8.07
adaptor-related protein complex 2, mu 1 subunit	NM_004068	7.95
interferon regulatory factor 4	NM_002460	7.90
ribosomal protein S19	NM_001022	7.90
interferon stimulated gene (20kD)	NM_002201	7.88
dolichyl-diphosphooligosaccharide-protein glycosyltransferase	D28643	7.80
actin, beta	X00351	7.72
major histocompatibility complex, class II, DR alpha	M60333	7.69
major histocompatibility complex, class I, B	L42024	7.62
major histocompatibility complex, class II, DR beta 5	AJ287586	7.57
actin, gamma 1	AL567820	7.49
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9)	NM_004542	7.48
HLA-B associated transcript 1	NM_004640	7.47
ribosomal protein L18a	AC004692	7.36
H2A histone family, member O	AJ313324	7.31
gb:U82824.1 /DEF=Homo sapiens HLA class I heavy chain (HLA-Cw*1701) mRNA, complete cds. /FEA=CDS /GEN=HLA-Cw*1701 /PROD=HLA class I heavy chain /DB_XREF U82824	U82824	7.31
major histocompatibility complex, class II, DR alpha	M60334	7.27
ribosomal protein S15	NM_001018	7.26
eukaryotic translation initiation factor 3, subunit 8 (110kD)	NM_003752	7.25
U8 snRNA-associated Sm-like protein LSm7	NM_016198	7.25
eukaryotic translation elongation factor 2	NM_001961	7.22
membrane-spanning 4-domains, subfamily A, member 1	X12530	7.13
eukaryotic translation initiation factor 3, subunit 8 (110kD)	AA678705	7.11
spermidine synthase	NM_003132	7.09
polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)	BG335629	7.09
ribosomal protein S2	NM_002952	7.03
chemokine (C-X-C motif), receptor 4 (lusin)	AF348491	7.03
similar to HYPOTHETICAL 34.0 KDA PROTEIN ZK795.3 IN CHROMOSOME IV	BC003829	7.00
RNA, U2 small nuclear	BE747342	6.98
origin recognition complex, subunit 5-like (yeast)	AF081459	6.94
DEAD-box protein abstrakt	NM_016222	6.93
immunoglobulin heavy constant mu	BC001872	6.90
DKFZP564M182 protein	AK000822	6.88
nuclear RNA helicase, DECD variant of DEAD box family	NM_005804	6.86
stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein)	BE866580	6.80
immunoglobulin heavy constant mu	BG340548	6.76
muS homolog 6 (E. coli)	D89846	6.75
tubulin, beta, 4	AL565749	6.75
major histocompatibility complex, class II, DR beta 1	U65585	6.73
T cell receptor beta locus	M15564	6.72
major histocompatibility complex, class II, DP alpha 1	M27487	6.62
ribonucleotide reductase M1 polypeptide	AI692874	6.62
signal sequence receptor, delta (translocon-associated protein delta)	NM_006280	6.59
seryl-RNA synthetase	NM_006513	6.55
eukaryotic translation initiation factor 2, subunit 3 (gamma, 52kD)	NM_001415	6.54
protein phosphatase 1, regulatory subunit 7	BF718769	6.52
ancient ubiquitous protein 1	NM_012103	6.52
glutathione S-transferase pi	NM_000852	6.49
polymerase (RNA) III (DNA directed) polypeptide K (12.3 kDa)	NM_016310	6.42
ubiquinol-cytochrome c reductase core protein 1	NM_003385	6.42
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	AB033605	6.41
splicing factor, arginine/serine-rich 5	AW084582	6.40

SKB1 homolog (S. pombe)	NM_006109	6.32
cell death-regulatory protein GRIM19	NM_015965	6.20
U6 snRNA-associated Sm-like protein	NM_012321	6.18
eukaryotic translation initiation factor 3, subunit 2 (beta, 36kD)	U36764	6.17
mitochondrial ribosomal protein S2	NM_016034	6.16
D123 gene product	NM_006023	6.15
replication factor C (activator 1) 2 (40kD)	NM_02914	6.09
membrane-spanning 4-domains, subfamily A, member 1	BC02807	6.02
Rab acceptor 1 (prenylated)	NM_006423	5.97
eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)	BC001173	5.97
ubiquitin C	M2880	5.94
valosin-containing protein	AF100752	5.93
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1	AL080089	5.93
CD22 antigen	X52785	5.89
B lymphoid tyrosine kinase	NM_001715	5.88
mitochondrial ribosomal protein S34	NM_023936	5.88
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASH1)	NM_005004	5.86
mitochondrial ribosomal protein L9	AB049636	5.85
ribosomal protein S2	A183766	5.84
actin binding LIM protein 1	NM_006720	5.84
syntactin 16	AK026970	5.83
major histocompatibility complex, class II, DR beta 4	NM_021863	5.82
protein kinase, DNA-activated, catalytic polypeptide	U34994	5.81
ribosomal protein S28	AC005011	5.80
eukaryotic translation initiation factor 4A, isoform 1	NM_001416	5.76
ribosomal protein, large, P0	BC003655	5.75
NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)	AF155235	5.75
ATP citrate lyase	U18197	5.74
Consensus includes gb:BF9794.18 /FEA=EST /DB_XREF=gi:12346634 /DB_XREF=est:60228246F1 /CLONE=IMAGE-4373914 /UG=Hs.119122 ribosomal protein L13a	BF9794.18	5.74
glutaminyl-HRNA synthetase	NM_005051	5.74
Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mrna /DB_XREF=gi:5915893 /UG=H	AF005487	5.73
eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)	NM_003751	5.71
polymerase (RNA) II (DNA directed) polypeptide L (7.6kD)	BC005903	5.70
mitochondrial ribosomal protein S12	NM_021107	5.66
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit 1, isoform 2	NM_004889	5.65
putative human HLA class II associated protein I	BE560202	5.64
hypothetical protein	NM_016459	5.58
membrane component, chromosome 11, surface marker 1	BG258784	5.57
hypothetical protein PRO1847	AF119855	5.57
proteasome (prosome, macropain) subunit, beta type, 7	NM_002799	5.57
cell division cycle 2-like 2	AF067524	5.57
similar to RIKEN cDNA 231004OG17 gene	BF972185	5.57
putative transmembrane protein; homolog of yeast Golgi membrane protein Yif1p (Yip1p-interacting factor)	NM_020470	5.57
tyllophanyl-RNA synthetase	M61715	5.56
ribosomal protein, large, P0	NM_001002	5.56
transcription factor Dp-1	NM_007111	5.55
guanine nucleotide binding protein (G protein), beta polypeptide 1	AF141124	5.54
actin, gamma 1	AU145192	5.53
hypothetical protein R33729_1	AC005339	5.53
CDW52 antigen (CAMPATH-1 antigen)	NM_001803	5.52
ribosomal protein, large, P0	AIS53822	5.51
actin related protein 23 complex, subunit 2 (34 kD)	AF279893	5.50
guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1	NM_006098	5.49
eukaryotic translation initiation factor 4A, isoform 1	BC006210	5.47
ubiquitin C	AB009010	5.47
polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)	AW402635	5.40
hypothetical protein MGC4875	AL118502	5.39
ribosomal protein L10	NM_006013	5.35
v-yes-1 Yarneguchi sarcoma viral related oncogene homolog	M79321	5.33
calpain, small subunit 1	AD001527	5.31
lymphocyte-specific protein tyrosine kinase	NM_005356	5.31
anaphase promoting complex subunit 5	BC001081	5.29
ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast)	BG395660	5.29

kinesin 2 (60-70kD)	AA284075	5.29
hematological and neurological expressed 1	NM_018185	5.27
ribosomal protein, large, P0	BC005863	5.26
immunoglobulin heavy constant mu	S74639	5.25
proteasome (prosome, macropain) subunit, beta type, 8 (large multifunctional protease 7)	U17496	5.24
ribosomal protein S3	U14990	5.22
proteasome (prosome, macropain) 26S subunit, ATPase, 3	U34690	5.21
coronin, actin binding protein, 1A	AF241788	5.17
nuclear distribution gene C (A. nidulans) homolog	NM_000631	5.17
neutrophil cytosolic factor 4 (40kD)	M24668	5.16
gb:M24668.1 /DEF=Human Ig rearranged H-chain V-region mRNA (C-D-JH4), complete cds. /FEA=mRNA /GEN=IGH@_DB_XREF=gi:185198 /FL=gb:M24668.1	NM_006924	5.16
splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor)	M12879	5.11
Cw1 antigen	BC004422	5.09
arabinose homolog 2 (Drosophila)	D42040	5.09
bromodomain-containing 2	AW514210	5.08
major histocompatibility complex, class I, F	NM_003000	5.08
succinate dehydrogenase complex, subunit B, iron sulfur (lp)	NM_004551	5.07
NADH dehydrogenase (ubiquinone) Fe-S protein 3 (30kD) (NADH-coenzyme Q reductase)	BC003573	5.01
farnesyl-diphosphate farnesyltransferase 1	NM_003584	4.98
transgelin 2	NM_001861	4.98
cytochrome c oxidase subunit IV isoform 1	NM_019058	4.95
integrin beta 4 binding protein	AF022229	4.95
HIF-1 responsive RTP801	NM_018573	4.93
hypothetical protein PRO1068	NM_014453	4.90
putative breast adenocarcinoma marker (32kD)	U20489	4.88
protein tyrosine phosphatase, receptor type, O	NM_025205	4.86
hypothetical protein DKFZP434N185	BC005946	4.85
tubulin alpha 6	AA789278	4.85
ribosomal protein L13	AI589086	4.83
Lysosomal-associated multispanning membrane protein-5	M18468	4.83
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	NM_001004	4.82
ribosomal protein, large P2	AF104913	4.79
eukaryotic translation initiation factor 4 gamma, 1	NM_007108	4.78
transcription elongation factor 8 (SIII), polypeptide 2 (18kD, elongin B)	NM_002271	4.76
karyopherin (importin) beta 3	J02923	4.74
lymphocyte cytosolic protein 1 (L-plastin)	L19184	4.70
peroxiredoxin 1	NM_016292	4.70
heat shock protein 75	AF228990	4.70
HLA-G histocompatibility antigen, class I, G	AF118710	4.69
ribosomal protein S14	NM_003592	4.69
culin 1	BC002526	4.68
heat shock 70kD protein 4	AF082131	4.65
NADH dehydrogenase (ubiquinone) flavoprotein 1 (51kD)	NM_014145	4.63
HSPC274 protein	BE305165	4.60
Consensus includes gb:BE305165 /FEA=EST /DB_XREF=gi:9177184 /DB_XREF=est:601186685T1 /CLONE=IMAGE:2959580 /UG=Hs.100623 phospholipase C, beta 3, neigl	D55716	4.60
MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)	NM_003406	4.56
lysine 3-monooxygenase/hypophan 5-monooxygenase activation protein, zeta polypeptide	U10485	4.55
lymphoid-restricted membrane protein	BG389015	4.53
tumor protein D52	J04564	4.53
small nuclear ribonucleoprotein polypeptides B and B1	U38979	4.52
postmeiotic segregation increased 2-like 9	NM_002466	4.51
v-myb myeloblastosis viral oncogene homolog (avian)-like 2	AF043179	4.46
T cell receptor beta locus	AW150953	4.45
7-dehydrocholesterol reductase	AL523776	4.45
hypothetical protein FLJ20113	BC000733	4.44
eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD)	AB028869	4.41
baculoviral IAP repeat-containing 3 (survivin)	NM_006152	4.39
lymphoid-restricted membrane protein	M80281	4.39
APEX nuclease (multifunctional DNA repair enzyme)	NM_018694	4.39
Consensus includes gb:Z82202 /DEF=Human DNA sequence from clone RP1-34P24 on chromosome 22 Contains a pseudogene similar to ribosomal protein L35, ESTs, STs	AA112507	4.38
HSV1 binding protein	AL559122	4.38
U6 snRNA-associated Sm-like protein		
T cell receptor beta locus		

casein kinase 2, beta polypeptide	NM_001320	4.38
putative methyltransferase	NM_017528	4.37
VPS28 protein	NM_016208	4.35
lysosomal-associated membrane protein 1	NM_005561	4.33
bone marrow stromal cell antigen 2	NM_004335	4.32
ribosomal protein L13a	BF942308	4.32
transmembrane protein 4	BC001027	4.32
calreticulin	AD000092	4.30
nuclear RNA export factor 1	BC004904	4.30
polymerase (RNA) II (DNA directed) polypeptide E (25kD)	AI554759	4.28
nucleosome assembly protein 1-like 4	NM_005989	4.25
gb:AY014272.1 /DEF=Homo sapiens FKSG30 (FKSG30) mRNA, complete cds, /FEA=mRNA (GEN=FKSG30 /PROD=FKSG30 /DB_XREF=gi:12408251 /UG=Hs.315492 Homc	AY014272	4.25
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	NM_006773	4.24
KIA00818 gene product	AI768378	4.23
LIM domain only 2 (homobin-like 1)	NM_005574	4.23
non-POU-domain-containing, octamer-binding	BC003129	4.22
peptidylprolyl isomerase F (cyclophilin F)	NM_005729	4.19
zinc finger protein 207	BE871379	4.18
enopamil binding protein (sterol isomerase)	AV702405	4.17
unknown	NM_016209	4.17
H31 binding protein	NM_006118	4.16
acetylase 2, mitochondrial	NM_001098	4.16
H3 histone, family 3B (H3.3B)	NM_005324	4.15
C2f protein	U72514	4.14
ribosomal protein L13a	BC001675	4.14
replication protein A1 (70kD)	NM_002945	4.12
prolasease (prosome, macropain) 26S subunit, non-ATPase, 2	NM_002808	4.12
gb:AY014272.1 /DEF=Homo sapiens FKSG30 (FKSG30) mRNA, complete cds, /FEA=mRNA (GEN=FKSG30 /PROD=FKSG30 /DB_XREF=gi:12408251 /UG=Hs.315492 Homc	NM_001550	4.11
interferon-related developmental regulator 1	M50684	4.11
HLA-G histocompatibility antigen, class I, G	NM_016337	4.11
RNB6	U23946	4.10
RNA binding motif protein 5	NM_003145	4.09
signal sequence receptor, beta (translocon-associated protein beta)	NM_005273	4.09
guanine nucleotide binding protein (G protein), beta polypeptide 2	NM_001029	4.08
ribosomal protein S28	M12423	4.07
T cell receptor alpha locus	NM_012458	4.07
translocase of inner mitochondrial membrane 13 homolog B (yeast)	NM_004111	4.06
flap structure-specific endonuclease 1	NM_014294	4.06
translocating chain-associating membrane protein	BC003381	4.06
KIAA0217 protein	NM_012456	4.05
translocase of inner mitochondrial membrane 10 homolog (yeast)	NM_005919	4.04
MADS box transcription enhancer factor 2, polypeptide B (myocyte enhancer factor 2B)	BC004949	4.03
tubulin alpha 6	M31183	4.03
major histocompatibility complex, class I, E	AF217197	4.02
fuse-binding protein-interacting repressor	BC005147	4.02
FK506 binding protein 1A (12kD)	AI987323	4.01
ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle	AF321125	4.00
DNA replication factor	BG028844	4.00
HLA-B associated transcript 3	NM_005001	4.00
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5kD, B14.5a)	U74223	4.00
poly(rC) binding protein 1	NM_005724	3.98
tetraspan 3	AI972144	3.98
hydroxyacyl-Coenzyme A dehydrogenase/3-ketocyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	NM_000075	3.98
cyclin-dependent kinase 4	BE560461	3.98
solute carrier family 2 (facilitated glucose/fructose transporter), member 5	BG257762	3.98
hypothetical protein	M15565	3.96
T cell receptor alpha locus	NM_006409	3.95
actin related protein 2/3 complex, subunit 1A (41 kD)	L34703	3.94
T cell receptor alpha locus	AW001847	3.94
amyloid beta (A4) precursor-like protein 2	NM_006327	3.93
translocase of inner mitochondrial membrane 23 homolog (yeast)	AL535380	3.93
B-cell translocation gene 1, anti-proliferative	BC003049	3.92
PAL-1 mRNA-binding protein	U39945	3.92
adenyate kinase 2		

cytochrome c oxidase subunit VIc
 TAP binding protein (tapsin)
 ribosomal protein L27a
 protein phosphatase 1, catalytic subunit, alpha isoform
 Consensus includes gb:SG531983 /FEA=EST /DB_XREF=gi:13523521 /DB_XREF=est:602561007F1 /CLONE=IMAGE:4699176 /UG=Hs.108104 ubiquitin-conjugating enzyme BG531983
 Consensus includes gb:SG3334495 /FEA=EST /DB_XREF=gi:13140933 /DB_XREF=est:602461128F1 /CLONE=IMAGE:4577718 /UG=Hs.228201 ESTs, Moderately similar to f BG334495
 leucine rich repeat (in FLJ) interacting protein 1
 ribosomal protein, large, P1
 cold shock domain protein A
 Tu translation elongation factor, mitochondrial
 ribosomal protein S18
 KIAA0748 protein
 protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform
 ubiquitin-activating enzyme E1 (A1SST and BN75 temperature sensitivity complementing)
 mitochondrial ribosomal protein S16
 8-phosphogluconolactonase
 CCAAT/enhancer binding protein (C/EBP), gamma
 major histocompatibility complex, class I, E
 coatomer protein complex, subunit epsilon
 chloride intracellular channel 1
 hypothetical protein FLJ20340
 CD27-binding (Siva) protein
 postmeiotic segregation increased 2-like 11
 Consensus includes gb:X02189 /DEF=H.sapiens adenosine deaminase (ADA) gene 5 flanking region and exon 1 (and joined CDS) /FEA=CDS /DB_XREF=gi:283558 /UG=Hs.1 X02189
 hypothetical protein
 Consensus includes gb:BF530535 /FEA=EST /DB_XREF=gi:11617898 /DB_XREF=est:602071788F1 /CLONE=IMAGE:4214680 /UG=Hs.77665 KIAA0102 gene product /FL=g BF530535
 FK506 binding protein precursor
 CDW52 antigen (CAMPATH-1 antigen)
 tyrosine 3-monooxygenase/tyriophan 5-monooxygenase activation protein, epsilon polypeptide
 translocase of inner mitochondrial membrane 44 homolog (yeast)
 neural precursor cell expressed, developmentally down-regulated 8
 H1 histone family, member X
 KIAA0618 gene product
 heterogeneous nuclear ribonucleoprotein R
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 4
 proteasome (prosome, macropain) subunit, beta type, 3
 glutamyl-prolyl-IRNA synthetase
 GCN1 general control of amino-acid synthesis 1-like 1 (yeast)
 eukaryotic translation elongation factor 1 gamma
 thymopollitin
 Consensus includes gb:BE731738 /FEA=EST /DB_XREF=gi:10145730 /DB_XREF=est:601568154F1 /CLONE=IMAGE:3842844 /UG=Hs.182937 peptidylprolyl isomerase A (c BE731738
 cytochrome c oxidase subunit Vb
 mitochondrial carrier homolog 2
 tyrosine 3-monooxygenase/tyriophan 5-monooxygenase activation protein, zeta polypeptide
 interferon enhancer binding factor 3, 90kD
 enhancer of invasion 10
 RuvB-like 2 (E. coli)
 small nuclear ribonucleoprotein polypeptide A
 hypothetical protein FLJ22837
 myosin 10
 cytochrome P450, subfamily 51 pseudogene 2
 SNRPN upstream reading frame
 ribosomal protein L13a
 H2B histone family, member G
 stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein)
 leucine-rich repeat protein, neuronal 1
 nuclear distribution gene C (A.ndulans) homolog
 hydroxyacyl-Coenzyme A dehydrogenase, type II
 MCM5 mitochondrial chromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)
 PTD008 protein
 eukaryotic translation elongation factor 1 alpha 1
 adenine phosphoribosyltransferase
 GDP-mannose 4,6-dehydratase

AA382702 3.91
 AF029750 3.91
 NM_000990 3.91
 NM_002708 3.91
 Consensus BG531983 3.90
 Consensus BG3334495 3.90
 NM_004735 3.89
 NM_001003 3.89
 NM_003651 3.89
 NM_003321 3.89
 NM_022551 3.88
 AA522514 3.88
 NM_014225 3.85
 NM_003334 3.85
 NM_016065 3.84
 NM_012088 3.84
 NM_001806 3.83
 NM_005516 3.82
 NM_007263 3.81
 AF034607 3.80
 NM_017773 3.80
 AF033111 3.80
 U38980 3.79
 X02189 3.79
 AF670847 3.78
 BF530535 3.78
 NM_016594 3.77
 N90866 3.76
 U28936 3.75
 NM_006351 3.75
 NM_006156 3.75
 NM_006026 3.75
 NZ5665 3.74
 NM_005826 3.74
 U51007 3.74
 NM_002795 3.73
 A142677 3.72
 A1697055 3.71
 NM_001404 3.71
 AF113682 3.71
 BE731738 3.71
 NM_001862 3.70
 NM_014342 3.68
 U28964 3.68
 AF147209 3.68
 NM_021178 3.67
 NM_006666 3.65
 NM_004596 3.65
 NM_025165 3.65
 AA621862 3.64
 U40053 3.64
 NM_022804 3.63
 NM_012423 3.63
 NM_003522 3.63
 AL553320 3.63
 A1654857 3.63
 NM_006600 3.63
 NM_004493 3.62
 NM_006739 3.62
 NM_016145 3.61
 NM_001402 3.60
 AA927724 3.60
 NM_001500 3.59

PRP9 pre-mRNA processing factor 8 homolog (yeast)
hepatitis delta antigen-interacting protein A
Consensus includes gb:Z95126 /DEF=Human DNA sequence from PAC 30P20 on chromosome Xq21.1-Xq21.3. Contains set pseudogene, ESTs and STS /FEA=COS /DB_XR_Z95128
Williams-Beuren syndrome chromosome region 5
guanylate kinase 1
intercellular adhesion molecule 2
nuclear protein family A, member 3 (HACA small nuclear RNPs)
G protein-coupled receptor 18
major histocompatibility complex, class II, DQ alpha 1
ribosomal protein L15
ATP-binding cassette, sub-family F (GCN20), member 2
KIAA0618 gene product
BCL2-interacting killer (apoptosis-inducing)
adaptor-related protein complex 2, sigma 1 subunit
Consensus includes gb:BF695847 /FEA=EST /DB_XREF=gi:11981255 /DB_XREF=est:601852205F1 /CLONE=IMAGE:4076232 /UG=Hs.301917 YDD19 protein
ARPP2 actin-related protein 2 homolog (yeast)
structure specific recognition protein 1
myosin, light polypeptide 6, alkali, smooth muscle and non-muscle
delta sleep inducing peptide, immunoreactive
Consensus includes gb:A001784 /FEA=EST /DB_XREF=gi:3202255 /DB_XREF=est:041g06.s1 /CLONE=IMAGE:1619386 /UG=Hs.306332 ESTs, Highly similar to A42735 rib A1001784
mercaptopyruvate sulfurtransferase
GNAS complex locus
hypothetical protein FLJ10359
DKFZP434J1813 protein
lymphotxin beta (TNF superfamily, member 3)
Consensus includes gb:AJ011414.1 /DEF=Homo sapiens mRNA for plexin-B1 plasma membrane receptor, truncated splice variant (plexin-B1SEP gene). /FEA=mRNA /GEN=AJ011414
DKFZP586M1523 protein
cytochrome c oxidase subunit Vb
ribonuclease H1
methionine-tRNA synthetase
immunoglobulin lambda locus
proteasome (prosome, macropain) 26S subunit, non-ATPase, 3
myosin, light polypeptide 6, alkali, smooth muscle and non-muscle
insulin induced gene 1
ATP synthase, H+ transporting, mitochondrial F1 complex, beta polypeptide
bryoid subunit 70kD (Ku antigen)
Consensus includes gb:AK027225.1 /DEF=Homo sapiens cDNA: FLJ23572 fis, clone LNG12403. /FEA=mRNA /DB_XREF=gi:10440306 /UG=Hs.201593 Homo sapiens cDNA: AK027225
branched chain aminotransferase 1, cytosolic
FK506 binding protein 1A (12kD)
Huntingtin interacting protein B
chromatin-specific transcription elongation factor, 140 kDa subunit
similar to mouse Gli3 or D. melanogaster transcription factor iIB
zinc finger protein 313
POP7 (processing of precursor, S. cerevisiae) homolog
structure specific recognition protein 1
mitochondrial ribosomal protein L23
isopentenyl-diphosphate delta isomerase
Xq28, 2000bp sequence contig, ORF
glutathione peroxidase 1
hypothetical protein MGC5585
GNAS complex locus
cytosolic acyl coenzyme A thioester hydrolase
phosphoinositide-3-kinase, regulatory subunit, polypeptide 3 (p55, gamma)
A kinase (PKA) anchor protein 2
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG)
serine hydroxymethyltransferase 2 (mitochondrial)
malic enzyme 2, NAD(+)-dependent, mitochondrial
oxygenol binding protein-like 2
HSPC142 protein
RNA binding motif protein 10
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4
DNA segment on chromosome 19 (unique) 1177 expressed sequence
threonine-tRNA synthetase

NM_008445 3.59
NM_008848 3.59
AF257135 3.58
BC006249 3.57
NM_008873 3.57
NM_018648 3.57
AF261135 3.55
BG397856 3.54
AF279803 3.54
NM_005692 3.54
AA514822 3.53
NM_001197 3.53
NM_021375 3.53
BF695847 3.52
NM_005722 3.51
NM_003146 3.51
AA419227 3.51
AL110191 3.50
NM_001784 3.49
NM_021126 3.45
AF064082 3.45
NM_018072 3.45
BG168666 3.45
NM_002341 3.44
NM_001414 3.44
BF063898 3.43
BC006229 3.41
NM_002936 3.41
NM_004990 3.40
XG3006 3.40
NM_002809 3.40
BE734356 3.39
BE300521 3.37
NM_001686 3.37
NM_001469 3.36
NM_001725 3.35
NM_005604 3.35
BC001002 3.34
AF049103 3.33
NM_007192 3.33
NM_013242 3.33
AL031685 3.32
BC001430 3.32
BE795648 3.31
AI832239 3.31
NM_004508 3.30
BE076218 3.29
NM_000581 3.29
NM_024057 3.28
AF088184 3.27
NM_007274 3.27
BE622627 3.27
BE679367 3.27
NM_004546 3.27
AW180316 3.26
M55905 3.26
BC000296 3.25
BC006244 3.25
AL137421 3.25
A1744900 3.25
NM_006114 3.25
NM_003191 3.25

Consensus includes gb:X04802/DEF=Homo sapiens UBBP2 pseudogene for ubiquitin UBB /FEA=CDS/DB_XREF=gi:37582 /UG=Hs.247890 Homo sapiens UBBP2 pseudoge	X04802	3.25
glyogenin	NM_004130	3.24
GNAS complex locus	NM_000516	3.24
ribosomal protein L10	AL031276	3.24
macrophage erythroblast attachr	NM_005882	3.24
Ewing sarcoma breakpoint region 1	BC000527	3.24
isodizate dehydrogenase 3 (NAD+) gamma	NM_004135	3.23
Consensus includes gb:AL118510/DEF=Human DNA sequence from clone RPS-858M22 on chromosome 20 Contains a pseudogene similar to 40S ribosomal protein S10, ST	AL118510	3.23
paired immunoglobulin-like receptor beta	NM_013440	3.23
KIAA0729 protein	AW502434	3.22
serine palmitoyltransferase, long chain base subunit 2	U15555	3.22
endothelial differentiation-related factor 1	AB002282	3.22
kinesin-like 4	AC002301	3.21
ubiquitin B	X04801	3.21
sialyltransferase 9 (CMP-NeuAc: lactosylceramide alpha-2,3-sialyltransferase; GM3 synthase)	NM_003896	3.21
eukaryotic translation elongation factor 1 alpha 1	NM_001403	3.20
chromosome 12 open reading frame 8	NM_006817	3.20
alanine-RNA synthetase	NM_001605	3.20
hypothetical protein FLJ11305	NM_018388	3.20
NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NM_002496	3.19
N-acylaminoacyl-peptide hydrolase	NM_001640	3.19
peptidylprolyl isomerase E (cyclophilin E)	AF042386	3.18
branched chain aminotransferase 1, cytosolic	AB022662	3.18
zinc finger protein 163 (RING finger, C2HC4 type)	BC000832	3.18
amino acid transporter 2	NM_018976	3.17
chromosome 14 open reading frame 3	NM_012111	3.17
SKI-interacting protein	NM_012111	3.17
interferon regulatory factor 3	AV689564	3.16
polyglutamine binding protein 1	NM_001571	3.16
ubiquitin specific protease 7 (herpes virus-associated)	AB041836	3.16
glutamic-pyruvate transaminase (alanine aminotransferase)	NM_003470	3.15
CD48 antigen (B-cell membrane protein)	NM_005309	3.15
golgi associated, gamma adenylin ear containing, ARF binding protein 2	NM_001778	3.15
oxidase (cytochrome c) assembly 1-like	BC000284	3.15
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	BC001669	3.15
RAN binding protein 1	AI122754	3.14
mitochondrial solute carrier	NM_002882	3.14
flightless I homolog (Drosophila)	BE677761	3.14
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD	AI830227	3.14
Consensus includes gb:BG537190 /FEA=EST/DB_XREF=gi:13528922 /DB_XREF=est:602565589F1 /CLONE=IMAGE:4690079 /UG=Hs.111334 ferritin, light polypeptide	NM_001695	3.14
CD27-binding (Sive) protein	BG537190	3.13
FK506 binding protein precursor	NM_006427	3.13
succinate dehydrogenase complex, subunit A, flavoprotein (Fp)	NM_016594	3.13
chromosome condensation 1	NM_004168	3.12
PRKC, apoptosis, W11, regulator	NM_001269	3.11
villin 2 (ezrin)	AI336206	3.11
glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1)	J05021	3.11
splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)	BC000498	3.11
hypothetical protein FLJ10969	BC000451	3.11
interleukin 21 receptor	NM_018292	3.10
G8 protein	NM_021798	3.10
tyrosine 3-monoxygenase/tryptophan 5-monoxygenase activation protein, theta polypeptide	NM_016947	3.10
isopentenyl-diphosphate delta isomerase	NM_006826	3.10
transforming growth factor, beta 1	BC005247	3.10
nicotinamide nucleotide transhydrogenase	BC000125	3.10
homeo box A1	NM_012343	3.10
CD37 antigen	AC004079	3.10
G1 to S phase transition 1	NM_001774	3.10
ribosomal protein S8	NM_002094	3.09
glycoprotein, synaptic 2	NM_001012	3.09
suppression of tumorigenicity 14 (colon carcinoma, multipase, epithin)	NM_004668	3.09
synaptophysin-like protein	NM_021878	3.09
proteasome (prosome, macropain) 26S subunit, non-ATPase, 9	AI768845	3.08
	NM_002813	3.08

Fanconi anemia, complementation group A	AV083279	3.08
zinc finger protein 259	NM_003904	3.07
translin	AA35302	3.07
WD repeat domain 1	AB010427	3.07
ubiquitin-conjugating enzyme E2M (UBC12 homolog, yeast)	NM_003969	3.07
CCAAT/enhancer binding protein (C/EBP), beta	AL564683	3.07
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 1	NM_004939	3.06
NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NM_002496	3.06
immunoglobulin lambda joining 3	AV698647	3.06
H2A histone family, member E	NM_021066	3.05
gb:AF008937.1 DEF-Homo sapiens syntaxin-16C mRNA, complete cds. /FEA=mRNA /PROD=syntaxin 16 /FL=gb:AF008937	gb:AF008937.1	3.05
block of proliferation 1	BC491842	3.05
interleukin enhancer binding factor 3, 90kD	AF141870	3.05
SMC4 structural maintenance of chromosomes 4-like 1 (yeast)	NM_009496	3.05
hypothetical protein FLJ12973	NM_024908	3.05
HLA-B associated transcript 3	BC003133	3.05
gb:Z25433.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405744 /FL=gb:Z25433	Z25433	3.04
inhibitor of DNA binding 1, dominant negative helix-loop-helix protein	D13889	3.03
Consensus includes gb:AA680302 /FEA=EST /DB_XREF=gi:2656270 /DB_XREF=IMAGE:969193 /UG=Hs.181125 immunoglobulin lambda locus	AA680302	3.03
transcriptional regulator protein	NM_013260	3.03
COP9 homolog	NM_008710	3.03
chromodomain protein, Y chromosome-like	AF081258	3.03
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	J04984	3.03
ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)	BE262760	3.03
B-cell associated protein	NM_007273	3.03
glutamic-oxaloacetic transaminase 2, mitochondrial (aspartate aminotransferase 2)	NM_002080	3.02
villin 2 (ezrin)	AA670344	3.02
GTP cyclohydrolase I feedback regulatory protein	NM_005256	3.02
eukaryotic translation elongation factor 1 gamma	AF119850	3.01
mitochondrial ribosomal protein L11	NM_016050	3.01
CD79A antigen (immunoglobulin-associated alpha)	NM_001783	3.01
ras homolog gene family, member A	BC001360	3.00
KIAA1007 protein	BC000779	3.00
presenilin 1 (Alzheimer disease 3)	NM_007318	2.99
KIAA0974 protein	BE551340	2.99
immunoglobulin lambda locus	M87790	2.99
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	NM_002716	2.98
solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	BF340083	2.98
solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2	NM_002394	2.98
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	M65254	2.98
inactive progesterone receptor, 23 kD	BC003005	2.97
Spogen's syndrome/deridroma autoantigen 1	NM_008396	2.97
HLA-B associated transcript 3	NM_004639	2.97
triophosphate isomerase 1	BF116254	2.97
B-cell CLL/lymphoma 6 (zinc finger protein 51)	S67779	2.96
major histocompatibility complex, class II, DR beta 5	NM_002125	2.96
nuclear and coiled-body phosphoprotein 1	NM_004741	2.96
hypothetical protein FLJ10534	NM_018128	2.96
Consensus includes gb:BF308250 /FEA=EST /DB_XREF=gi:11255432 /DB_XREF=est:601890308F1 /CLONE=IMAGE:4131673 /UG=Hs.315103 ESTs. Moderately similar to C	BF308250	2.95
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; KI)	BC002684	2.94
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 56/58kD, isoform 2	NM_001693	2.94
manic fringe homolog (Drosophila)	A1736965	2.94
sperm associated antigen 9	NM_003971	2.93
epoietin antagonizing transcription factor	AF063208	2.93
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 21kD	BC005876	2.93
HSPC003 protein	NM_014017	2.93
nuclear prelamin A recognition factor	NM_012336	2.93
ribosomal protein L5	NM_000869	2.92
sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)	AV695711	2.92
soc-2 suppressor of clear homolog (C. elegans)	NM_007373	2.91
adenosine kinase	U90339	2.91
mitochondrial ribosomal protein S7	NM_015971	2.91
CD38 antigen (p45)	NM_001775	2.91

major histocompatibility complex, class I, C
 Consensus includes gb:AL353681 /DEF=Human DNA sequence from clone RP1-158P9 on chromosome 1. Contains a putative novel gene, a laminin receptor 1 (67kD, ribosom
 hypothetical protein FLJ10439
 adenosine deaminase
 membrane component, chromosome 11, surface marker 1
 chromosome 20 open reading frame 1
 arachidonate 5-lipoxygenase-activating protein
 fusion, derived from t(12;18) malignant liposarcoma
 casein kinase 2, alpha 1 polypeptide
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8 (19kD, PGIV)
 HLA-G histocompatibility antigen, class I, G
 potassium channel, subfamily K, member 12
 HSPC037 protein
 SH3 domain binding glutamic acid-rich protein like 3
 sphingosine-1-phosphatase
 vesicle-associated membrane protein 1 (synaptobrevin 1)
 ubiquitin-like 5
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily f, member 1
 thyroid receptor interacting protein 15
 Consensus includes gb:AK000185.1 /DEF=Homo sapiens cDNA FLJ20178 fts, clone COL09990. /FEA=nrRNA.DB_XREF=gi:7020104 /UG=Hs.306389 Homo sapiens cDNA FI
 pericentriolar material 1
 CDC16 cell division cycle 16 homolog (S. cerevisiae)
 acute carrier family 1 (neutral amino acid transporter), member 5
 dynein, cytoplasmic, heavy polypeptide 1
 GDP-mannose pyrophosphorylase A
 ATX1 antioxidant protein 1 homolog (yeast)
 fragile X mental retardation, autosomal homolog 1
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1
 H2A histone family, member X
 Consensus includes gb:AA468687 /FEA=EST.DB_XREF=gi:2216118 /DB_XREF=estab3803.s1 /CLONE=IMAGE:843101 /UG=Hs.284235 ESTs
 spectrin SH3 domain binding protein 1
 CCR4-NOT transcription complex, subunit 8
 proteasome (prosome, macropain) subunit, beta type, 4
 CGI-47 protein
 phosphoserine aminotransferase
 nuclear matrix protein NMP200 related to splicing factor PRP19
 transforming growth factor, beta 2
 replication factor C (activator 1) 5 (36.5kD)
 RNA binding motif protein 4
 splicing factor 3a, subunit 3, 60kD
 cytoskeleton-associated protein 1
 SH3-domain, GRB2-like, endophilin B2
 neural precursor cell expressed, developmentally down-regulated 5
 serine/threonine kinase 15
 DnaJ (Hsp40) homolog, subfamily C, member 8
 msp kinase phosphatase-like protein MK-STYX
 CD70B antigen (immunoglobulin-associated beta)
 uridine monophosphate kinase
 hypothetical protein PRO1847
 chromosome 20 open reading frame 80
 interferon stimulated gene (20kD)
 similar to S. cerevisiae RER1
 adaptor-related protein complex 2, sigma 1 subunit
 eukaryotic translation initiation factor 4E-like 3
 eukaryotic translation initiation factor 5A
 Consensus includes gb:A591100 /FEA=EST.DB_XREF=gi:4600148 /DB_XREF=est1w01cd3.x1 /CLONE=IMAGE:2267044 /UG=Hs.113368 neuroendocrine secretory protein
 gb:M24669.1 /DEF=Human 17 rearranged H-chain V-region mRNA (C-D-JH6), complete cds. /FEA=mrRNA /GEN=IGH@ /DB_XREF=gi:185200 /FL=gb:M24669.1
 Breakpoint cluster region protein, uterine leiomyoma, 1: barrier to autointegration factor
 cytoskeleton-associated protein 1
 RAD23 homolog A (S. cerevisiae)
 hypothetical protein FLJ20859
 membrane interacting protein of RGS16
 polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)

AK024838 2.90
 BC006351 2.90
 NM_000022 2.89
 NM_005898 2.89
 AF098158 2.89
 NM_001629 2.89
 S75762 2.89
 A1181318 2.89
 NM_014222 2.89
 M90666 2.88
 NM_022055 2.88
 BC003186 2.88
 NM_031286 2.88
 NM_030791 2.88
 NM_016830 2.87
 NM_024292 2.87
 AF231056 2.87
 NM_004236 2.87
 AK000185 2.86
 NM_006197 2.86
 NM_003903 2.86
 AF105230 2.86
 AB002323 2.86
 NM_013335 2.85
 NM_004045 2.85
 A1990766 2.85
 AK021419 2.85
 NM_002105 2.85
 AA48687 2.85
 BF873013 2.84
 A1769416 2.84
 NM_002796 2.84
 NM_016014 2.84
 A1889380 2.83
 NM_014502 2.83
 NM_003238 2.83
 NM_007370 2.83
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 NM_006802 2.82
 NM_001281 2.81
 NM_020145 2.81
 NM_004404 2.81
 NM_003600 2.80
 NM_014280 2.80
 NM_016086 2.80
 NM_000626 2.80
 BC002806 2.79
 AF119855 2.79
 AL441988 2.79
 U88964 2.79
 NM_007033 2.78
 NM_004089 2.78
 AF047695 2.77
 NM_001870 2.76
 A1591100 2.76
 M24669 2.76
 AF044773 2.76
 BC005869 2.75
 BF572938 2.75
 NM_022734 2.75
 NM_016641 2.75
 AA679988 2.75

TNF-induced protein	BC005352	2.75
HLA-G histocompatibility antigen, class I, G	M90685	2.74
deoxyguanosine kinase	NM_001929	2.74
KIAA1228 protein	AJ708767	2.74
cut-like 1, CCAAT displacement protein (Drosophila)	NM_001913	2.74
endonuclease G	NM_004435	2.74
CREBBP/EP300 inhibitory protein 1	AF274951	2.74
damage-specific DNA binding protein 1 (127kD)	L40326	2.74
RAN binding protein 7	AL137335	2.73
sorting nexin 5	NM_014426	2.73
hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	U04627	2.73
transportin-SR	NM_012470	2.72
24-dehydrocholesterol reductase	NM_014762	2.72
microtubule-associated proteins 1A/1B light chain 3	AF183417	2.72
GDP dissociation inhibitor 2	NM_001494	2.72
bleomycin hydrolase	NM_000386	2.72
DNA segment, single copy probe LNS-CAILNS-CALI (deleted in polyposis)	A4814140	2.72
major histocompatibility complex, class II, DM alpha	X76775	2.72
adaptor-related protein complex 2, sigma 1 subunit	BC006337	2.72
UDP glycosyltransferase 2 family, polypeptide B15	NM_001076	2.72
Consensus includes gb:AW582267 /FEA=EST /DB_XREF=gi:7257316 /DB_XREF=gi:7257316 /DB_XREF=est:QVQ-ST0215-060100-083-c06 /UG=Hs.30695.1 Human DNA sequence from clone RP11	AW582267	2.72
MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	A859865	2.71
laminin receptor 1 (67kD, ribosomal protein SA)	AL136306	2.71
KIAA0095 gene product	NM_014669	2.71
serine/threonine kinase 24 (STE20 homolog, yeast)	AA586774	2.71
RAB5A, member RAS oncogene family	NM_004162	2.71
nuclear factor (erythroid-derived 2)-like 1	A361227	2.70
solute carrier family 7 (cationic amino acid transporter, y+ system), member 5	AB018009	2.70
src family associated phosphoprotein 2	AC003999	2.70
RD RNA-binding protein	L03411	2.70
lymphocyte-specific protein tyrosine kinase	U07236	2.70
cytochrome c-1	NM_01916	2.70
core histone macroH2A2.2	NM_018649	2.70
deoxyhypusine synthase	U26266	2.70
hypothetical protein MGC10433	NM_024321	2.69
Consensus includes gb:BE988833 /FEA=EST /DB_XREF=gi:10579538 /DB_XREF=est:60164986 /F1 /CLONE=IMAGE:3933782 /UG=Hs.324648 Homo sapiens cDNA FLJ137	BE988833	2.69
immunoglobulin lambda joining 3	X57812	2.69
zinc finger protein 143 (clone pHZ-1)	AJ277546	2.69
3'-phosphoadenosine 5'-phosphosulfate synthase 1	AF033026	2.69
Fanconi anemia, complementation group G	AC004472	2.69
RAB4, member RAS oncogene family	NM_004578	2.69
postmeiotic segregation increased 2-like 3	D38437	2.68
gb:NM_016592.1 /DEF=Homo sapiens neuroendocrine secretory protein 55 (NESP55), mRNA. /FEA=mRNA /GEN=NESP55 /PROD=neuroendocrine secretory protein 55 /DB_XREF=gi:10579538 /DB_XREF=est:60164986 /F1 /CLONE=IMAGE:3933782 /UG=Hs.324648 Homo sapiens cDNA FLJ137	NM_016592	2.68
adenine phosphoribosyltransferase	NM_000485	2.68
hypothetical protein R32184_1	BC001648	2.68
deoxyhypusine synthase	NM_001930	2.68
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	AF081496	2.67
voltage-dependent anion channel 1	AL151918	2.67
Rho GDP dissociation inhibitor (GDI) beta	NM_001175	2.67
ribosomal protein L24	NM_000986	2.67
gb:NM_018604.1 /DEF=Homo sapiens hypothetical protein PRO1741 (PRO1741), mRNA. /FEA=mRNA /GEN=PRO1741 /PROD=hypothetical protein PRO1741 /DB_XREF=gi:10579538 /DB_XREF=est:60164986 /F1 /CLONE=IMAGE:1127826 /UG=Hs.132390 zinc finger protein 38 (KOX 18)	NM_018604	2.67
Consensus includes gb:AA653300 /FEA=EST /DB_XREF=gi:2569471 /DB_XREF=est:ag65c10.s1 /CLONE=IMAGE:1127826 /UG=Hs.132390 zinc finger protein 38 (KOX 18)	AA653300	2.66
muscleblind-like (Drosophila)	N31913	2.66
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	NM_002734	2.66
high-mobility group 20B	AF288679	2.66
anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2)	BE869563	2.66
H2A histone family, member Y	AF044286	2.66
casein kinase 2, beta polypeptide	M30448	2.65
zinc finger protein 43 (HITF8)	NM_003423	2.65
hypothetical protein MGC10715	AL049650	2.65
mesenchymal stem cell protein DSC43	A1992095	2.65
proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)	NM_002800	2.65
adenylyate kinase 2	W02312	2.65

mesenchymal stem cell protein DSC92	NM_016645	2.85
transcription factor CP2	U03494	2.85
polymerase (RNA) II (DNA directed) polypeptide 1 (14.5kD)	AL037557	2.84
polymerase (RNA) II (DNA directed) polypeptide D	NM_004805	2.84
RNA-binding protein (autoantigenic)	NM_016732	2.83
hepatocellular carcinoma-associated antigen 66	NM_016428	2.83
ADP-ribosylation factor 1	AA580004	2.83
v-rel-1 murine leukemia viral oncogene homolog 1	NM_002880	2.83
triosephosphate isomerase 1	NM_000385	2.83
seb4D	AL109955	2.83
polymerase (RNA) III (DNA directed) (62kD)	BC004424	2.83
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), member D	AL566172	2.82
thioredoxin domain-containing	NM_030755	2.82
FK506 binding protein 1A (12kD)	NM_000801	2.81
cyclochrome b-245, alpha polypeptide	NM_000101	2.81
eukaryotic translation initiation factor 4E binding protein 1	AB044548	2.81
aldo-keto reductase family 1, member A1 (aldohyde reductase)	NM_008066	2.81
archelin 1	NM_001655	2.81
death associated protein 3	U16321	2.81
RuvB-like 1 (E. coli)	NM_003707	2.81
RNA binding motif protein, X chromosome	AA52524	2.80
fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)	NM_023107	2.80
mutS homolog 5 (E. coli)	AF034759	2.80
hydroxyamide S-acyltransferase (E2 component of pyruvate dehydrogenase complex)	J03866	2.80
ribosomal protein L44	NM_021029	2.80
high-mobility group (nonhistone chromosomal) protein 14	NM_004965	2.80
Ras-GTPase activating protein SH3 domain-binding protein 2	NM_012297	2.59
60S ribosomal protein L30 isolog	NM_016304	2.59
AD021 protein	NM_016613	2.59
uncoupling protein 2 (mitochondrial, proton carrier)	U82819	2.59
60S acidic ribosomal protein PO	NM_016183	2.58
C-terminal binding protein 1	NM_001328	2.58
ATP synthase, H ⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1	NM_005174	2.58
insulin induced gene 1	NM_005542	2.58
chloride intracellular channel 4	AF109196	2.58
ribosomal protein S25	AA888388	2.58
SEC13-like 1 (S. cerevisiae)	NM_030673	2.58
survival of motor neuron protein interacting protein 1	AB037701	2.58
CGI-150 protein	AF061730	2.57
hypothetical protein FLJ12287 similar to semaphorins	NM_022367	2.57
proline synthetase co-transcribed (bacterial homolog)	NM_007198	2.57
ribosomal protein S8	BC000524	2.57
breast carcinoma amplified sequence 4	NM_017843	2.57
phosphoglucanate dehydrogenase	NM_002631	2.57
transferrin receptor (p80, CD71)	NM_003234	2.57
neural precursor cell expressed, developmentally down-regulated 5	AI191427	2.57
enkephase promoting complex subunit 5	BF667271	2.57
MAP kinase-interacting serine/threonine kinase 1	BC002755	2.57
calnexin	L18887	2.56
gila maturation factor, gamma	NM_004877	2.56
ATPase, Na ⁺ /K ⁺ transporting, beta 3 polypeptide	U51478	2.56
sorting nexin 1	AF065484	2.55
Rho GDP dissociation inhibitor (GDI) alpha	BC005851	2.55
histidyl-RNA synthetase	NM_002109	2.55
platelet/endothelial cell adhesion molecule (CD31 antigen)	M37760	2.55
XUAA0123 protein	BF570122	2.55
ret finger protein	NM_018150	2.55
interleukin 21 receptor	AF230394	2.55
ARP3 actin-related protein 3 homolog (yeast)	AF269133	2.55
ADP-ribosylation factor 4	NM_005721	2.54
heterogeneous nuclear ribonucleoprotein F	AL537042	2.54
phorbol-12-myristate-13-acetate-induced protein 1	AI591354	2.54
	AI857639	2.53

cAMP responsive element modulator	NM_001881	2.53
ubiquitin-conjugating enzyme E2 variant 1	U39361	2.53
immunoglobulin lambda locus	AF043586	2.53
chromosome condensation-related SMC-associated protein 1	AK022511	2.53
hypothetical protein MGC4368	NM_024510	2.53
fer (lpafes related) tyrosine kinase (phosphoprotein NCP94)	NM_005246	2.53
chromatin assembly factor 1, subunit A (p150)	NM_005483	2.52
estrogen-related receptor alpha	L39487	2.52
hypothetical protein MGC5242	NM_024033	2.52
glyceraldehyde-3-phosphate dehydrogenase	AK026525	2.52
hypothetical protein FLJ10702	NM_018184	2.51
valyl-tRNA synthetase 2	NM_006295	2.51
heat shock 90kD protein 1, alpha	AI062933	2.51
villin 2 (ezrin)	BF663141	2.51
desitin (actin depolymerizing factor)	NM_021130	2.50
peptidylprolyl isomerase A (cyclophilin A)	NM_015640	2.50
PAL-1 mRNA-binding protein	NM_000284	2.50
pyruvate dehydrogenase (lipoamide) alpha 1	NM_016007	2.50
gb:NM_016007.1/DEF=Homo sapiens CGI-59 protein (LOC51625), mRNA, /FEA=mRNA /GEN=LOC51625 /PROD=CGI-59 protein /DB_XREF=gi:7706297 /UG=Hs.279887 C	NM_016007	2.49
protein phosphatase 1, regulatory (inhibitor) subunit 2	NM_008241	2.49
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 17 (72kD)	U59321	2.49
glycyl-tRNA synthetase	D30658	2.49
putative translation initiation factor	BF246436	2.49
putative translation initiation factor	Z97353	2.48
ribosomal protein L15	NM_024006	2.48
hypothetical protein IMAGE3455200	NM_017916	2.48
hypothetical protein FLJ20643	NM_024579	2.48
hypothetical protein FLJ23221	BC000924	2.48
NBR2	AL537707	2.48
putative translation initiation factor	U54829	2.47
regulator of G-protein signalling 16	NM_006503	2.47
proteasome (prosome, macropain) 26S subunit, ATPase, 4	AA167775	2.47
NRAS-related gene	NM_003340	2.47
ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)	NM_030978	2.47
hypothetical protein similar to actin related protein 2/3 complex, subunit 5	AF189015	2.47
villin 2 (ezrin)	NM_002949	2.46
mitochondrial ribosomal protein L12	BC004334	2.46
ribosomal protein S10	BC289867	2.46
RAD21 homolog (S. pombe)	NM_001428	2.46
endolase 1, (alpha)	NM_021633	2.46
kelch-like protein C3IP1	AL136932	2.45
inhibitor of growth family, member 1	AF044076	2.45
putative cyclin G1 interacting protein	NM_006349	2.45
CD184 antigen, alalomonuch	BF669455	2.45
ubiquitin-cytochrome c reductase hinge protein	NM_006004	2.45
asparyl aminopeptidase	NM_012100	2.45
chaperonin containing TCP1, subunit 8 (theta)	NM_004599	2.43
superoxide dismutase 1, soluble (erythrocyte lateral sclerosis 1 (adult))	NM_002764	2.43
zinc finger protein	D38553	2.43
isocitrate dehydrogenase 3 (NAD+) beta	AI050286	2.43
steroid regulatory element binding transcription factor 2	NM_018455	2.43
phosphoribosyl pyrophosphate synthetase 1	BC000027	2.43
KIAA0074 protein	NM_001942	2.43
hypothetical protein MGC4022	AF116273	2.43
uncharacterized bone marrow protein BM039	NM_001518	2.43
integral type I protein	BC000649	2.42
Consensus includes gb:M21388.1 /DEF=Human unproductively rearranged Ig mu-chain mRNA V-region (VD), 5' end, clone mu-3A1A, /FEA=mRNA /GEN=IGHM /DB_XREF=g	BF343852	2.42
desmoglein 1	AA630314	2.42
BCL2-associated athanogene		
general transcription factor II, i		
ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1		
histone H2A.F/Z variant		
ribosomal protein S2		

Consensus includes gb:BE963164 /FEA=EST /DB_XREF=gi:11766582 /DB_XREF=est:601656973R1 /CLONE=IMAGE:3865650 /UG=Hs.2186 eukaryotic translation elongation BE963164 2.42
 IGF-II mRNA-binding protein 3 AU160004 2.42
 Consensus includes gb:AL121975 /DEF=Human DNA sequence from clone RP3-422B11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the P AL121975 2.42
 proteolipid protein 2 (colonic epithelium-enriched) BC005982 2.42
 peptidylprolyl isomerase A (cyclophilin A) NM_030666 2.41
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 NM_014402 2.41
 low molecular mass ubiquitinone-binding protein (9.5kD) NM_016576 2.41
 GUPR2 for guanosine monophosphate reductase isolog NM_007096 2.41
 cathrin, light polypeptide (Lca) U38964 2.40
 postmeiotic segregation increased 2-like 8 AV007161 2.40
 SRY (sex determining region Y)-box 2 N29889 2.40
 UBX domain-containing 2 A1628925 2.40
 KIAA1895 protein NM_020381 2.40
 candidate tumor suppressor protein NM_014183 2.40
 HSPC162 protein AA502912 2.39
 KIAA0906 protein NM_004526 2.39
 MCM2 minichromosome maintenance deficient 2, mitotin (S. cerevisiae) BC003574 2.39
 T-cell leukemia/lymphoma 1A NM_002573 2.39
 platelet-activating factor acetylhydrolase, isoform 1b, gamma subunit (29kD) NM_005769 2.39
 proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) AL518328 2.39
 basic leucine-zipper protein BZAP45 NM_020187 2.39
 DC12 protein BF342707 2.38
 pre-mRNA processing factor 31 homolog (yeast) AF074723 2.38
 RNA polymerase II transcriptional regulation mediator (Med6, S. cerevisiae, homolog of) NM_003712 2.38
 protein phosphatase 1, regulatory subunit 7 NM_003730 2.38
 ribonuclease 6 precursor NM_003932 2.38
 suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein) NM_003932 2.38
 zinc finger protein 195 AA284075 2.38
 karesin 2 (60-70kD) AF164622 2.38
 golgin-47 NM_001786 2.37
 cell division cycle 2, G1 to S and G2 to M BG034328 2.37
 transcription factor Op-2 (E2F dimerization partner 2) AF683882 2.37
 Rho-associated, coiled-coil containing protein kinase 1 gb:AF274954, 1 /DEF=Homo sapiens PNAS-29 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-29 /DB_XREF=gi:12751070 /UG=Hs.328791 Homo sapiens PNAS-29 mRNA AF274954 2.37
 HSPC002 protein NM_015362 2.37
 gb:NM_004682.1 /DEF=Homo sapiens PCA and SFRS1 interacting protein 2 (PSIP2), mRNA. /FEA=mRNA /GEN=PSIP2 /PROD=PCA and SFRS1 interacting protein 2 /DB_XF NM_004682 2.37
 hypothetical protein MGC2722 BF975273 2.36
 uncharacterized hypothalamus protein HT009 NM_018470 2.36
 lipok acid synthetase BF056778 2.36
 dUTP pyrophosphatase U90223 2.36
 mitochondrial ribosomal protein L17 AK026857 2.36
 ribosomal protein S15a NM_001019 2.36
 dynactin 1 (p150, glued homolog, Drosophila) NM_004082 2.35
 polymerase (RNA) II (DNA directed) polypeptide G NM_002696 2.35
 H2B histone family, member A BC000893 2.35
 major histocompatibility complex, class I, A A4573862 2.35
 activity-dependent neuroprotector NM_015339 2.35
 mitochondrial ribosomal protein S17 NM_015969 2.35
 Consensus includes gb:BE250348 /FEA=EST /DB_XREF=gi:9120458 /DB_XREF=est:600943342T1 /CLONE=IMAGE:2959811 /UG=Hs.326249 Homo sapiens mRNA; cDNA [BE250348 2.35
 FYTD domain-containing ion transport regulator 5 NM_014164 2.34
 actin related protein 2/3 complex, subunit 5 (16 kD) AL516350 2.34
 maltrin 3 A4577695 2.34
 general transcription factor II, i BC004472 2.34
 Wolf-Hirschhorn syndrome candidate 1 BE793789 2.34
 casein kinase 2, alpha prime polypeptide NM_001896 2.34
 nuclear receptor co-repressor 1 AA085748 2.34
 synovial sarcoma translocation, chromosome 18 NM_005637 2.33
 eukaryotic translation elongation factor 1 alpha 1 AL515273 2.33
 casein kinase 1, alpha 1 NM_001892 2.33
 methylcrotonoyl-Coenzyme A carboxylase 2 (beta) AB050049 2.33
 prostaglandin D2 synthase, hemopoietic NM_014485 2.33
 stem-loop (histone) binding protein NM_006527 2.33
 isocitrate dehydrogenase 3 (NAD+) alpha A1828060 2.33

hypothetical protein FLJ21617	NM_030897	2.33
ribosomal protein S12	AI799007	2.33
kinesin-like 6 (mitotic centromere-associated kinesin)	U63743	2.33
chromosome condensation 1	X06130	2.33
ribophorn II	BC003560	2.33
protein phosphatase 6, catalytic subunit	NM_027271	2.33
splicing factor, arginine/serine-rich 7 (35kD)	AA524053	2.33
Wolf-Hirschhorn syndrome candidate 1	AF083389	2.33
serum/glucocorticoid regulated kinase-like	NM_013257	2.33
guanine nucleotide binding protein (G protein), alpha 13	NM_006572	2.32
MYG1 protein	NM_021640	2.32
FtsJ homolog 2 (E. coli)	AK024635	2.32
accessory proteins BAP31/BAP29	NM_005745	2.32
TRAF family member-associated NFKB activator	NM_004180	2.32
cytochrome b5 outer mitochondrial membrane precursor	NM_030579	2.32
hypothetical protein MGC4504	NM_024111	2.32
pyridoxal (pyridoxine, vitamin B6) kinase	NM_003681	2.32
protease, serine, 22	NM_022119	2.31
ribosomal protein L4	A053886	2.31
zinc finger protein 225	AC074331	2.31
general transcription factor IIC, polypeptide 3 (102kD)	NM_012086	2.31
SMY3 suppressor of mit two 3 homolog 1 (yeast)	BG338532	2.31
hypothetical protein FLJ20436	AK022732	2.31
mitogen-activated protein kinase kinase 7	AF218074	2.31
D-dopa-chrome tautomerase	NM_001355	2.31
translocase of outer mitochondrial membrane 20 (yeast) homolog	NM_014755	2.31
KIAA0755 gene product	A141892	2.31
RAD23 homolog B (S. cerevisiae)	AL527385	2.31
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	NM_000351	2.31
HSPC053 protein	NM_014151	2.31
6-phosphogluconolactonase	NM_012088	2.31
mannosidase, alpha, class 1A, member 1	NM_005807	2.31
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki)	AA758755	2.31
adenylosuccinate lyase	NM_000028	2.30
alkylglycerone phosphate synthase	NM_003659	2.30
dendritic cell protein	BC001465	2.30
HBS1-like (S. cerevisiae)	NM_006360	2.30
hypothetical protein FLJ12806	NM_022831	2.30
amyloid beta (A4) precursor-like protein 2	BG427393	2.29
X-prolyl aminopeptidase (aminopeptidase P)-like	W07689	2.29
PK domain containing 16	NM_006523	2.29
ring finger protein 10	NM_022114	2.29
high-mobility group (nonhistone chromosomal) protein 1	NM_014868	2.29
WW domain binding protein 11	BE311760	2.29
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	AF118023	2.29
ATP-dependent interferon responsive	NM_001945	2.29
neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	NM_022371	2.28
peptidyl prolyl isomerase H (cyclophilin H)	AW072386	2.28
tryptophanyl-tRNA synthetase	NM_006347	2.28
major histocompatibility complex, class I, C	NM_004184	2.28
KIAA1228 protein	BC004489	2.28
postmeiotic segregation increased 2-like 1	BC001224	2.28
eukaryotic translation initiation factor 28, subunit 2 (beta, 39kD)	D38435	2.28
DKFZP586A011 protein	NM_014239	2.28
H3 histone family, member L	NM_015416	2.28
Finkel-Biskamp-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (for derived); ribosomal protein S30	NM_003537	2.27
hypothetical protein PRO1635	NM_001997	2.27
hypothetical protein MGC2718	NM_016589	2.27
seven in absentia homolog 2 (Drosophila)	NM_024067	2.27
protein tyrosine phosphatase, non-receptor type 22 (lymphoid)	U76248	2.27
nuclear transcription factor Y, gamma	NM_015967	2.27
hypothetical protein FLJ23209	U82296	2.27
	NM_024895	2.27

hypothetical protein FLJ12549	NM_024844	2.27
BRCA1 associated protein	AF035620	2.27
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)	AW188131	2.26
glyoxylate reductase/hydroxypyruvate reductase	NM_012203	2.26
collagen, type VI, alpha 1	M20776	2.26
protein kinase domains containing protein similar to phosphoprotein C8FW	NM_021158	2.26
progesterin induced protein	AF006010	2.25
HSPC160 protein	NM_014182	2.25
zinc finger protein 9 (a cellular retroviral nucleic acid binding protein)	NM_003418	2.25
H2A histone family, member Y	AF044286	2.25
polymerase (DNA-directed), delta 4	NM_021173	2.25
divalent cation tolerant protein CUTA	AF230924	2.25
hypothetical protein AF140225	NM_030799	2.25
immunoglobulin lambda locus	AF072445	2.25
isocitrate dehydrogenase 3 (NAD+) beta	AF023265	2.25
gb:D13287.1/DEF+Human mRNA for alternative splicing product of glucocerebrosidase pseudogene, complete cds. /FEA=mRNA (PROD=glucocerebrosidase /DB_XREF=gi4 D13287	AB049944	2.25
mitochondrial ribosomal protein S11	BF976260	2.24
transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)	A1701949	2.24
heterogeneous nuclear ribonucleoprotein K	AA684258	2.24
heterogeneous nuclear ribonucleoprotein C (C1/C2)	U93240	2.24
karyopherin alpha 4 (importin alpha 3)	NM_014303	2.24
pescedillo homolog 1, containing BRCT domain (zebrafish)	NM_003315	2.24
DnaJ (Hsp40) homolog, subfamily C, member 7	BC003584	2.24
dihydrofolate reductase	U49188	2.24
tumor differentially expressed 1	NM_021968	2.24
H4 histone family, member E	NM_001628	2.24
aldo-keto reductase family 1, member B1 (aldose reductase)	BC006301	2.24
anaphase promoting complex subunit 5	BF689284	2.24
proliferation-associated ZC4, 38kD	NM_017654	2.24
hypothetical protein FLJ20512	BC000425	2.24
protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	AF263279	2.23
CD164 antigen, sialomucin	NM_005863	2.23
neuroepithelial cell transforming gene 1	BC000436	2.23
endosulfine alpha	NM_006414	2.23
ribonuclease P (38kD)	NM_012394	2.23
profilin 2	NM_001233	2.23
gb:BC001233.1/DEF+Homo sapiens. Similar to KIAA0092 gene product, clone MGC-4896, mRNA, complete cds. /FEA=mRNA (PROD=Similar to KIAA0092 gene product /DB_XREF=gi4 BC001233	NM_007317	2.23
capping protein (actin filament) muscle Z-line, beta	NM_004930	2.23
ribosomal protein S10	NM_001014	2.23
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 31kD	BC004443	2.23
KIAA0310 gene product	SC001404	2.23
ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-like 1	AL359763	2.23
claf lip and palate associated transmembrane protein 1	BC004865	2.23
3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (soluble)	NM_002130	2.23
statelin, RNA binding protein (Drosophila)	AL136601	2.23
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16kD	NM_001694	2.22
bone morphogenetic protein 7 (osteogenic protein 1)	M60316	2.22
Consensus includes gb:AW054828 /FEA=EST /DB_XREF=gi:5920529 /DB_XREF=est:ws60c12.x1 /CLONE=IMAGE:2501590 /UG=Hs.58220 Homo sapiens cDNA; FLJ23005	NM_002802	2.22
proteasome (prosome, macropain) 26S subunit, ATPase, 1	NM_015904	2.22
translation initiation factor IF2	NM_002676	2.22
phosphomannomutase 1	AB009358	2.22
mitogen-activated protein kinase kinase 7	NM_024050	2.22
hypothetical protein MGC2594	AF030514	2.21
small inducible cytokine subfamily B (Cys-X-Cys), member 11	D13989	2.21
Rho GDP dissociation inhibitor (GDI) alpha	AL136897	2.21
hypothetical protein FLJ11301	BC001149	2.21
serologically defined colon cancer antigen 16	NM_012392	2.21
PEF protein with a long N-terminal hydrophobic domain (pellin)	BC001886	2.21
ribonucleotide reductase M2 polypeptide	AK025270	2.21
hypothetical protein FLJ21617	NM_004309	2.21
Rho GDP dissociation inhibitor (GDI) alpha	AF180476	2.21
CCR4-NOT transcription complex, subunit 8		

intercellular adhesion molecule 3	NM_002162	2.21
KIAA00592 gene product	AI123527	2.21
HSPC144 protein	NM_014174	2.20
tubulin, alpha 1 (testis specific)	K03460	2.20
postmeiotic segregation increased 2-like 8	AB017004	2.20
mitochondrial ribosomal protein S18B	BC005373	2.20
5-enolmethylidene-4-carboxamide ribonucleotide formyltransferase/IMP cyclohydrolase	D89976	2.20
Cluster incl. AA131335:zb0801.s1 Homo sapiens cDNA, 3' end / clone=IMAGE-587097 / clone_end=3 / gb=AA131335 / gi=1692833 / ug=Hs.28166 / len=552	AA131335	2.20
ribosomal protein S21	NM_001024	2.20
poly(C) binding protein 2	NM_005016	2.20
NOL1R protein	AW248552	2.20
neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	NM_002265	2.20
programmed cell death 8 (apoptosis-inducing factor)	NM_004208	2.20
gb:M24670.1 / DEEF+Human Ig rearranged H-chain V-region mRNA (C-D-JH6), complete cds. / FEA=mRNA / GEN=IGH@ / DB_XREF=gi:165202 / FL=gb:M24670.1	M24670	2.20
methylene adenosyltransferase II, alpha	NM_005911	2.19
TIA1 cytosolic granule-associated RNA binding protein	BF692742	2.19
apoptosis related protein APR-3	NM_016085	2.19
ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)	NM_016085	2.19
FUS interacting protein (serine-arginine rich) 2	M32721	2.18
ubiquitination factor E48 (UFD2 homolog, yeast)	AU130523	2.19
CDC37 cell division cycle 37 homolog (S. cerevisiae)	AB028839	2.19
hypothetical protein FLJ11200	U63131	2.19
Consensus includes gb:AF052117.1 / DEEF+Homo sapiens clone 23809 mRNA sequence. / FEA=mRNA / DB_XREF=gi:3360424 / UG=Hs.6932 / Homo sapiens clone 23809 mRNA	NM_018359	2.19
polymerase (RNA) III (DNA directed) (62kD)	AF052117	2.19
nucleic acid diphosphate linked moiety X-type motif 1	U93867	2.19
hypothetical protein DKFp781H221	NM_002452	2.18
ATPase, H+ transporting, lysosomal (vacuolar proton pump), member J	NM_017601	2.18
kinesin-like 6 (mitotic centromere-associated kinesin)	BC003564	2.18
gb:NM_016402.1 / DEEF+Homo sapiens hypothetical protein (HSPC140), mRNA. / FEA=mRNA / GEN=HSPC140 / PROD=hypothetical protein / DB_XREF=gi:7705472 / UG=Hs.27	AY026505	2.18
hypothetical protein FLJ10479	NM_016402	2.18
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 35	NM_018106	2.18
hypothetical protein FLJ20257	NM_021831	2.18
KIAA1513 protein	NM_019606	2.18
CDC8 cell division cycle 8 homolog (S. cerevisiae)	NM_017542	2.18
H2A histone family, member C	NM_001254	2.17
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	NM_003509	2.17
inositol polyphosphate-5-phosphatase, TSKD	NM_003530	2.17
thousand and one amino acid protein kinase	AU160695	2.17
ribosomal protein S3A	A039084	2.17
hypothetical protein FLJ11301	AW007163	2.17
CASP2 and RIPK1 domain containing adaptor with death domain	A925635	2.17
karyopherin (importin) beta 1	AL136897	2.17
protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform	U79115	2.17
FK506 binding protein 2 (13kD)	AC004941	2.17
voltage-dependent anion channel 3	AA974416	2.16
hypothetical gene CG018	NM_004470	2.16
glutaryl-prolyl-4-aminopeptidase	U80943	2.16
similar to yeast BET3 (S. cerevisiae)	AL048785	2.16
ferritin, heavy polypeptide pseudogene 1	NM_004446	2.16
homolog of yeast MCM10; hypothetical protein PRO2249	AF041432	2.16
KIAA1049 protein	J04755	2.16
ribosomal protein L35a	NM_018518	2.16
hypothetical protein FLJ21610	AF322111	2.16
hypothetical protein FLJ20312	BE988801	2.16
synthrophin, beta 1 (dytrophin-associated protein A1, 59kD, basic component 1)	NM_022751	2.16
mitochondrial ribosomal protein L32	NM_017761	2.16
UDP-Gal:betaGalNAc beta 1,4-galactosyltransferase, polypeptide 3	NM_021021	2.15
chromosome 11 open reading frame2	NM_014180	2.15
nucleolar and coiled-body phosphoprotein 1	AF038661	2.15
hypothetical protein	D21262	2.15
hypothetical protein FLJ12298	NM_013265	2.15
3-hydroxy-3-methylglutaryl-Coenzyme A reductase	NM_015684	2.15
	AK022360	2.15
	AL518627	2.15

gb:NM_017445.1 /DEF=Homo sapiens H2B histone family, member S (H2BFS), mRNA, /FEA=CDS /GEN=H2BFS /PROD=H2B histone family, member S /DB_XREF=gi:11036	2.15
dykeritosis congenita 1, dykeritin	2.15
NAD kinase	2.15
chondroitin sulfate proteoglycan 6 (barnacan)	2.15
calcium-regulated heat-stable protein (24kD)	2.15
goli transport complex 1 (90 kDa subunit)	2.15
KIAA1858 protein	2.14
potassium channel, subfamily K, member 1 (TWIK-1)	2.14
solute carrier family 8 (neurotransmitter transporter, GABA), member 13	2.14
RAB7, member RAS oncogene family	2.14
hypothetical protein	2.14
postmeiotic segregation increased 2-like 8	2.14
actin, gamma 1	2.14
lymphocyte adaptor protein	2.14
carboxylate (N-acetylglucosamine 6-O) sulfotransferase 6	2.14
jumping translocation breakpoint	2.14
cytochrome b5 outer mitochondrial membrane precursor	2.14
ATP-binding cassette, sub-family F (GCN20), member 1	2.14
SHC (Src homology 2 domain-containing) transforming protein 1	2.14
RAD51 homolog C (S. cerevisiae)	2.13
hypothetical protein	2.13
Consensus includes gb:574774.1 /DEF=p59Yn(T)=OKT3-induced calcium influx regulator human, Jurkat J6 T cell line, mRNA, Partial, 1605 nt, /FEA=CDS /PROD=tyrosine kin, S74774	2.13
hematopoietic cell-specific Lyn substrate 1	2.12
RAP1, GTP-GDP dissociation stimulator 1	2.12
pne-B-cell colony-enhancing factor	2.12
2,5'-oligoadenylate synthetase 1 (40-46 kD)	2.12
hypothetical protein FLJ10803	2.12
adaptor protein with pleckstrin homology and src homology 2 domains	2.12
hypothetical protein	2.12
prostate tumor over expressed gene 1	2.12
chromodomain protein, Y chromosome-like	2.12
postmeiotic segregation increased 2-like 9	2.12
mitotic control protein dis3 homolog	2.12
guanine nucleotide binding protein (G protein), beta 5	2.12
FK506 binding protein 1A (12kD)	2.12
eukaryotic transition initiation factor 3, subunit 5 (epsilon, 47kD)	2.12
cystathyl-RNA synthetase	2.11
pyruvate-5-carboxylate synthetase (glutamate gamma-semialdehyde synthetase)	2.11
survival of motor neuron protein interacting protein 1	2.11
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4 (9kD, MLRQ)	2.11
HSPC023 protein	2.11
H2B histone family, member R	2.11
PI-3-kinase-related kinase SMG-1	2.11
myosin light chain 1 slow 8	2.11
RNA binding motif protein 5A	2.11
immediate early protein	2.11
nuclear cap binding protein subunit 2, 20kD	2.11
phosphatidylinositol glycan, class F	2.11
BCL2-associated athanogene 5	2.10
chromosome 2 open reading frame 3	2.10
KIAA1393 protein	2.10
prostatic binding protein	2.10
H2B histone family, member K	2.10
acid phosphatase 1, soluble	2.10
Consensus includes gb:BF786470 /FEA=EST /DB_XREF=gi:12101524 /DB_XREF=est:602259269F1 /CLONE=IMAGE:4342999 /UG=Hs.284275 Homo sapiens PAK2 mRNA, BF786470	2.10
FK506 binding protein 5	2.10
hypothetical protein FLJ20989	2.10
similar to S. pombe dim1+	2.10
eINOS interacting protein	2.10
ribosomal protein S4, X-linked	2.10
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	2.10
Wilms' tumour 1-associating protein	2.10
KIAA0623 protein	2.09

small glutamine-rich teratocopeptide repeat (TPR)-containing huntingtin interacting protein 2	NM_003021	2.09
DKFZP564F1123 protein	AB022435	2.09
acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A thiolase)	U346504	2.09
heat shock factor binding protein 1	NM_000019	2.09
gb:225425.1 /DEF=H.sapiens protein-serine/threonine kinase (DB_XREF=gi:405728 /FL=gb:225425 EC5IT	NM_001537	2.09
clathrin, heavy polypeptide (Hc)	NM_016581	2.09
nucleotide binding protein 2 (MinD homolog, E. coli)	NM_004859	2.09
deoxy-diphosphoglycerate-protein glycosyltransferase	NM_012225	2.09
NADH dehydrogenase (ubiquinone) Fe-S protein 6 (13kD) [NADH-coenzyme Q reductase]	BC002594	2.09
KIAA0164 gene product	NM_004553	2.09
TAF12 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 20 kD	BE963370	2.09
Inhibitor of DNA binding 2, dominant negative helix-loop-helix protein	D50544	2.09
CGI-120 protein	NM_002166	2.06
alpha-actinin-2-associated LIM protein	NM_016057	2.08
DKFZP434D156 protein	AF002280	2.08
tumor necrosis factor receptor superfamily, member 7	W93787	2.08
up-regulated by BCG-CWS	NM_001242	2.07
crystallin, beta B2	NM_022154	2.07
CGI-11 protein	NM_000496	2.07
apolipoprotein M	AF112204	2.07
MCM5 minichromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)	AF161434	2.07
DKFZP586J0619 protein	AA807528	2.07
H2B histone family, member B	BFS13430	2.07
polymerase (RNA) II (DNA directed) polypeptide H	BC002842	2.07
HSPC134 protein	U37689	2.07
RAP2A, member of RAS oncogene family	NM_014169	2.07
DNA-damage-inducible transcript 3	BE669921	2.07
tumor protein D52	BC003637	2.07
KIAA1104 protein	BE974088	2.07
prostate cancer overexpressed gene 1	NM_014968	2.06
splicing factor, arginine/serine-rich 2, interacting protein	NM_003627	2.06
phosphodiesterase 8D, cGMP-specific, rod, delta	NM_004719	2.06
postmeiotic segregation increased 2-like 2	NM_002801	2.06
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3	AA161026	2.06
nuclear receptor subfamily 3, group C, member 1	NM_001689	2.06
hypothetical protein DKFZ434E2216	U01351	2.06
ubiquitin specific protease 24	AC004475	2.06
chemokine-like factor 1	AB028880	2.06
high-mobility group (nonhistone chromosomal) protein 1	BF034561	2.06
eukaryotic translation elongation factor 1 alpha 1	NM_016951	2.06
general transcription factor IIH, polypeptide 1 (62kD subunit)	AC007277	2.06
cytochrome c oxidase subunit VIII	BE786672	2.05
SMC4 structural maintenance of chromosomes 4-like 1 (yeast)	NM_005316	2.05
gb:NM_000970.1 /DEF=Homo sapiens ribosomal protein L6 (RPL6), mRNA, /FEA=mRNA /GEN=RPL6 /PROD=ribosomal protein L6 /DB_XREF=gi:4506656 /UG=Hs.174131 r	NM_004074	2.05
protein phosphatase 1, regulatory (inhibitor) subunit 15A	AL136877	2.05
phenylalanine-RNA synthetase-like	NM_014330	2.05
apolipoprotein M	NM_004461	2.05
KIAA1856 protein	NM_019101	2.05
trifoli factor 1 (breast cancer, estrogen-inducible sequence expressed in)	AK026933	2.05
DKFZP566Q084 protein	NM_003225	2.05
gb:U72518.1 /DEF=Human desitin-2 pseudogene mRNA, complete cds, /FEA=mRNA /PROD=desitin-2 /DB_XREF=gi:1673523 /UG=Hs.195299 Human desitin-2 pseudogene	NM_015510	2.05
maleate dehydrogenase 1, NAD (soluble)	U72518	2.05
adaptor-related protein complex 3, delta 1 subunit	NM_005817	2.05
glyoxylate reductase/hydroxypyruvate reductase	AF130042	2.04
Consensus includes gb:AW408767 /FEA=EST /DB_XREF=gi:692782 /DB_XREF=gi:692782 /UG=Hs.293381 ESTs, Modera	AK026752	2.04
heterogeneous nuclear ribonucleoprotein A3	AW408767	2.04
synixin 7	N32035	2.04
origin recognition complex, subunit 5-like (yeast)	NM_002553	2.04
prolactosome (prolactin, macrophage) subunit, alpha type 6	BC002879	2.04
tumor differentially expressed 1	BC006088	2.04

synaptic nuclei expressed gene 2	NM_015180	2.03
stearyl-CoA desaturase (delta-9-desaturase)	BC005807	2.03
peptide deformylase-like protein	NM_022341	2.03
WD repeat domain 6	NM_018031	2.03
actophilin-1 interacting protein 1; actinin receptor interacting protein 1	AC004990	2.03
catalase	AY028632	2.03
ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)	NM_001419	2.03
SWAP-70 protein	AB014540	2.03
serologically defined colon cancer antigen 16	BC001149	2.03
transformation/transcription domain-associated protein	AC004893	2.03
tumor necrosis factor receptor superfamily, member 17	NM_001192	2.03
LIM and SH3 protein 1	NM_006148	2.03
nuclear cap binding protein subunit 1, 80kD	BC001450	2.03
KIAA1014 protein	AB023231	2.03
Wolff-Hirschhorn syndrome candidate 1	BF111870	2.03
DnaJ (Hsp40) homolog, subfamily A, member 2	AF011793	2.03
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g	AL050277	2.03
ornithine decarboxylase 1	NM_002539	2.03
defender against cell death 1	NM_001344	2.02
H2A histone family, member Z	NM_002106	2.02
Or-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	U77413	2.02
protease, serine, 25	AF184911	2.02
O-6-methylguanine-DNA methyltransferase	NM_002412	2.02
keratin, hair, acidic.3A	NM_004138	2.02
capillary renal cell carcinoma (translocation-associated)	BC004913	2.02
guanine nucleotide binding protein (G protein), beta 5	AF300650	2.02
calumenin	AF257659	2.02
GCM1 general control of amino-acid synthesis 1-like 1 (yeast)	D86973	2.02
cyclin D3	NM_001760	2.02
actin, gamma 1	AL515810	2.02
ribosomal protein L7a	NM_000972	2.02
B-cell translocation gene 1, anti-proliferative	NM_001731	2.01
KIAA1608 protein	NM_024820	2.01
stearyl-CoA desaturase (delta-9-desaturase)	AA678241	2.01
Consensus includes gb:AL137162 /DEF=Human DNA sequence from clone RP5-943L14 on chromosome 20. Contains ESTs, STSs and GSSs. Contains a novel gene and the vitamin A responsive; cytoskeleton related	AL137162	2.01
KIAA0225 protein	N92494	2.01
chromosome 11 open reading frame 10	AW208115	2.01
eukaryotic translation elongation factor 2	NM_014206	2.01
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	A004246	2.01
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9 (39kD)	AC004382	2.01
proteasome (prosome, macropain) activator subunit 2 (PA28 beta)	AF050641	2.01
fibroblast growth factor (acidic) intracellular binding protein	NM_002818	2.01
hippocalcin-like 1	NM_004214	2.01
ras homolog gene family, member H	NM_002149	2.00
non-canonical ubiquitin conjugating enzyme 1	NM_004310	2.00
similar to hypothetical protein MNCb-2386	NM_016021	2.00
hypothetical protein FLJ20346	AA421957	2.00
metallothionein 2A	NM_017872	2.00
coated vesicle membrane protein	NM_005953	2.00
NOL1R protein	NM_006815	2.00
melic enzyme 2, NAD(+)-dependent, mitochondrial	NM_018044	2.00
PRP4/STX/WD splicing factor	BC000147	2.00
putative dimethyladenosine transferase	U82756	2.00
tumor necrosis factor receptor superfamily, member 6	W87688	2.00
delta sleep inducing peptide, immunoreactor	X63493	2.00
hypothetical protein	AA1780381	2.00
survival of motor neuron protein interacting protein 1	BE314601	2.00
ubiquitin A-52 residue ribosomal protein fusion product 1	AB037702	1.99
olfactory receptor, family 1, subfamily K, member 1	NM_015956	1.99
forkhead box M1	AF348700	1.99
	NM_016835	1.99
	NM_021953	1.99

ankyrin 1, erythrocytic
 RAE1 RNA export 1 homolog (S. pombe)
 hypothetical protein FLJ20244
 NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)
 nudix (nucleoside diphosphate linked moiety X)-type motif 4
 heterogeneous nuclear ribonucleoprotein C (C1/C2)
 H2A histone family, member Y
 sperm associated antigen 9
 myeloid cell leukemia sequence 1 (BCL2-related)
 CDC8 cell division cycle 6 homolog (S. cerevisiae)
 cell division cycle 2, G1 to S and G2 to M
 calcium/calmodulin-dependent protein kinase IG
 Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)
 BCL2-associated athanogene
 ubiquinol-cytochrome c reductase core protein II
 Link guanine nucleotide exchange factor II
 core-binding factor, beta subunit
 DAZ associated protein 2
 hypothetical protein FLJ22382
 general transcription factor IIH, polypeptide 4 (52KD subunit)
 tyrosine kinase 2
 phosphoprotein enriched in astrocytes 15
 p47
 hypothetical protein MGC2855
 protein-L-isopartate (D-aspartate) O-methyltransferase
 secretory carrier membrane protein 3
 hypothetical nuclear factor SBB122
 DnaJ (Hsp40) homolog, subfamily B, member 6
 UDP-GlcNAc:beta-Gal beta-1,3-N-acetylglucosaminyltransferase 1
 protein kinase C and casein kinase substrate in neurons 2
 H2A histone family, member A
 hypothetical protein FLJ12860
 DnaJ (Hsp40) homolog, subfamily C, member 8
 gp.BC000285.1 /DEF-Homo sapiens, clone MGC:3182, mRNA, complete cds. /FEA=mrna /PROD=Unknown (protein for MGC:3182) /DB_XREF=gi:12653008 /UG=Hs.31541
 ubiquitin-conjugating enzyme HBUCE1
 diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)
 RNA-binding protein regulatory subunit
 sumnocalcin 2
 prohibitin
 proteasome (prosome, macropain) subunit, beta type, 10
 hypothetical protein FLJ20552
 KIAA1033 protein
 ATP-binding cassette, sub-family E (ABCF), member 1
 protein transport protein SEC81 alpha subunit isoform 1
 estrogen-related receptor alpha
 PTD016 protein
 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 1
 pyruvate dehydrogenase (lipoamide) beta
 hypothetical protein FLJ20707
 signal sequence receptor, alpha (translocin-associated protein alpha)
 solute carrier family 16 (monocarboxylic acid transporters), member 1
 myosin, light polypeptide, regulatory, non-sarcomeric (20KD)
 asparaginoglucosaminidase
 hypothetical protein FLJ10326
 protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform
 ATPase, Ca++ transporting, plasma membrane 1
 Sec23 homolog A (S. cerevisiae)
 eukaryotic translation elongation factor 1 beta 2
 replication factor C (activator 1) 4 (37KD)
 proteasome (prosome, macropain) subunit, alpha type, 5
 peroxiredoxin 2
 NOL1R protein

AF005213 1.99
 U65943 1.99
 NM_017722 1.99
 NM_005008 1.98
 NM_019094 1.98
 NM_004500 1.98
 NM_004893 1.98
 AB011088 1.98
 BF594446 1.98
 U77949 1.98
 D88357 1.98
 AW016039 1.97
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 NM_004323 1.97
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 AF264326 1.97
 AL534321 1.97
 NM_025045 1.97
 NM_001517 1.97
 NM_003331 1.97
 BC002426 1.97
 NM_016143 1.97
 NM_024339 1.97
 M83008 1.97
 NM_005698 1.97
 NM_020395 1.96
 AF080569 1.96
 NM_006577 1.96
 NM_007229 1.96
 NM_021062 1.96
 NM_025152 1.96
 AA83895 1.96
 BC000265 1.96
 W74577 1.96
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 AK35826 1.96
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 NM_002801 1.96
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 AK001657 1.96
 A002002 1.96
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 M65541 1.95
 NM_006364 1.95
 NM_001959 1.94
 NM_002916 1.94
 NM_002780 1.94
 NM_005809 1.94
 A0284845 1.94

lpyrRNA synthetase	AF285758	1.94
glutaredoxin (thioltransferase)	AF162789	1.94
aldolase A, fructose-bisphosphate	AK026577	1.94
JAK binding protein	AB005043	1.94
hypothetical protein FLJ11856	AK021918	1.94
glutathione S-transferase M1	NM_000561	1.94
KIAA0211 gene product	NM_014630	1.94
X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD)	J04977	1.94
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5 (13kD, B13)	NM_005000	1.93
diaphanous homolog 1 (Drosophila)	AF051782	1.93
hypothetical protein FLJ20317	NM_017859	1.93
galactosidase, beta 1	NM_000404	1.93
mannose phosphate isomerase	NM_002435	1.93
hypothetical protein FLJ10374	NM_018074	1.93
glutathione peroxidase 4 (phospholipid hydroperoxidase)	NM_002085	1.93
phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase	NM_000819	1.93
arsenate resistance protein ARS2	BE646076	1.93
Rho-associated, coiled-coil containing protein kinase 2	D87831	1.93
H2B histone family, member H	NM_003523	1.93
cell division cycle 27	NM_001256	1.93
vesicle docking protein p115	NM_003715	1.92
KIAA0174 gene product	NM_014761	1.92
KIAA0603 gene product	NM_014832	1.92
Consensus includes gb:AV715578 /FEA=EST /DB_XREF=est:AV715578 /CLONE=DCBB/G09 /UG=Hs.21056 Homo sapiens cDNA: FLJ21366 fis. cdc AV715578	NM_014832	1.92
Consensus includes gb:AL121916 /DEF=Human DNA sequence from clone RP11-189G13 on chromosome 20. Contains an RPL7A (60S ribosomal protein L7A) (SURF3) pseud	NM_014832	1.92
chromosome 1 open reading frame 16	NM_014837	1.92
cystathionine-beta-synthase	BE613178	1.92
hypothetical protein	NM_016618	1.92
slit3-associated polypeptide, 30kD	BF247098	1.92
polymerase I and transcript release factor	AF312393	1.92
RBP1-like protein	AA887480	1.92
signal peptidase 12kDa	NM_014041	1.92
aldolase A, fructose-bisphosphate	NM_000034	1.92
Ops-interacting protein 2	AL050353	1.92
staufen, RNA binding protein (Drosophila)	NM_004602	1.92
glutathione synthetase	NM_000178	1.92
interferon regulatory factor 7	NM_004030	1.92
N-myc downstream-regulated gene 3	BE300252	1.92
heterogeneous nuclear ribonucleoprotein A3	AA528233	1.92
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	BC003621	1.92
cyclin-dependent kinase 5	NM_004935	1.91
CD47 antigen (Rn-related antigen, integrin-associated signal transducer)	Z25521	1.91
microtubule-associated protein, RP/EB family, member 1	NM_004935	1.91
gb:AF660511.1 /DEF=Homo sapiens clone 016b10 My016 protein mRNA, complete cds. /FEA=mrna /PROD=My016 protein /DB_XREF=gi:12001971 /UG=Hs.181634 Homo	NM_012325	1.91
SET translocation (myeloid leukemia-associated)	M93851	1.91
ATPase, H+ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9	NM_005765	1.91
PWP2 periodic tryptophan protein homolog (yeast)	U56085	1.91
ubiquitin carboxyl-terminal esterase L1 (ubiquitin thioesterase)	NM_004181	1.91
POU domain, class 2, associating factor 1	NM_008235	1.91
McKusick-Kaufman syndrome	NM_018848	1.91
DKFZp568D133 protein	AL050050	1.91
guanine monophosphate synthetase	NM_003875	1.91
cysteinyl-RNA synthetase	NM_001751	1.91
DEAD/H (Asp-Glu-Ala-AspHis) box polypeptide 30	NM_014966	1.91
synplektin; Huntinglin interacting protein 1	Y10931	1.91
I-complex 1	BC000665	1.91
dopamine receptor D2	M30625	1.91
hypothetical protein FLJ20764	NM_017955	1.91
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16kD	M62762	1.91
gamma-tubulin complex protein 2	NM_006659	1.91
Consensus includes gb:AE653730 /FEA=EST /DB_XREF=gi:4737709 /DB_XREF=est:wb36112.x1 /CLONE=IMAGE:2307791 /UG=Hs.279009 matrix Gla protein	AE653730	1.91
CHK2 checkpoint homolog (S. pombe)	BC004207	1.91
transcription factor-like 4	BF056105	1.90

24

glutamate receptor, ionotropic, AMPA 3	NM_000828	1.87
coatomer protein complex, subunit alpha	U24105	1.87
GABA(A) receptor-associated protein	NM_007278	1.86
synaptic nuclei-expressed gene 1b	AF043290	1.86
H1 histone family, member 4	AL353759	1.86
chromosome 11 open reading frame 9	AC004770	1.86
S-phase kinase-associated protein 2 (p45)	BC001441	1.86
transcription factor Dp-1	R60866	1.86
similar to Bos laurus P14 protein	NM_030877	1.86
hypothetical protein FLJ13962	NM_024862	1.86
jumping translocation breakpoint	AF151058	1.86
leptin receptor overlapping transcript-like 1	AF161461	1.86
actin, beta	X00351	1.86
BTG3 associated nuclear protein	NM_017869	1.86
acid phosphatase 1, soluble	BE872974	1.86
nucleophosmin/nucleoplamin 3	NM_006993	1.86
osteoclast stimulating factor 1	NM_012383	1.86
RAB6A, member RAS oncogene family	NM_002869	1.86
epilepsy, progressive myoclonus type 2, Lafora disease (laforin)	BC005286	1.86
SVV/SHF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	D28156	1.86
actin, beta	AA809056	1.86
basic leucine zipper transcription factor, ATF-like	AC007182	1.86
aquaporin 7	NM_001170	1.86
ribosomal protein L10	AW057781	1.86
Consensus includes gb:18Q249599 /FEA=EST /DB_XREF=gi:12759426 /DB_XREF=est:602319692F1 /CLONE=IMAGE:4414895 /UG=Hs.110771 Homo sapiens cDNA: FLJ211	BG249599	1.85
ataxia telangiectasia and Rad3 related	U49844	1.85
chrate synthase	BC000105	1.85
actin, gamma 1	NM_001614	1.85
CGI-96 protein	AL022316	1.85
dynactin 3 (p22)	NM_007234	1.85
poly(A) binding protein, cytoplasmic 1	A1734929	1.85
hypothetical protein FLJ10036	NM_017875	1.85
PAP-1 binding protein	NM_031288	1.85
talin 1	NM_006289	1.85
ring finger protein 14	NM_004290	1.85
hypothetical protein FLJ12542	NM_024899	1.85
KIAA0542 gene product	AA521233	1.85
homolog of yeast RNase MRP/RNase P protein Pop5	NM_015918	1.85
cleavage and polyadenylation specific factor 1, 160kD subunit	NM_013291	1.84
poly(A) binding protein, nuclear 1	NM_004643	1.84
nucleoside phosphorylase	NM_000270	1.84
coristatin	AB000263	1.84
glucose regulated protein, S6kD	DB3485	1.84
procollagen (type III) N-endopeptidase	NM_002768	1.84
RNA binding motif protein 14	NM_006328	1.84
casein kinase 1, delta	NM_001893	1.84
H4 histone family, member D	NM_003541	1.84
suppressor of Ty, 5 homolog (S. cerevisiae)	NM_003169	1.84
MMS19-like (MET18 homolog, S. cerevisiae)	NM_022382	1.84
CDC14 cell division cycle 14 homolog A (S. cerevisiae)	AF064103	1.84
Consensus includes gb:AL354872 /DEF=Human DNA sequence from clone RP11-42015 on chromosome 1. Contains ESTs, STSs, GSSs and a CpG island. Contains the CTH	AL354872	1.84
KIAA0699 protein	BC002327	1.84
eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)	BC002513	1.84
protein kinase, cAMP-dependent, catalytic, beta	NM_002731	1.84
chaperonin containing TCP1, subunit 6A (zeta 1)	NM_001762	1.84
ligalin	NM_006893	1.84
RAP2B, member of RAS oncogene family	NM_002886	1.84
quinoid dihydropteridine reductase	BC000376	1.84
mitochondrial ribosomal protein L24	NM_024540	1.84
transmembrane protein 4	NM_014255	1.83
reticulon 2	AF004222	1.83
homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	AF217990	1.83
phorbol-12-myristate-13-acetate-induced protein 1	NM_021127	1.83

actin, beta	NM_001101	1.83
tyrosine kinase, non-receptor, 1	NM_003985	1.83
hypothetical protein FLJ13812	BC002449	1.83
acidic protein rich in leucines	AV712577	1.83
ribonuclease H1	BG534527	1.83
immunoglobulin heavy constant mu	BG340670	1.83
SET translocation (myeloid leukemia-associated)	D4S198	1.83
proteasome (prosome, macropain) subunit, alpha type, 1	NM_002786	1.83
hypothetical protein FLJ20897	AI613383	1.83
carbonyl reductase	NM_016286	1.83
DKFZP566C0424 protein	BG171020	1.82
non-metastatic cells 2, protein (NM23B) expressed in	NM_002512	1.82
Consensus includes gb:N92498/FEA-EST/DB_XREF=gi:1264807/DB_XREF=est:zb28a04.s1 CLONE=IMAGE:304878 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fa, N92498	NM_007006	1.82
cleavage and polyadenylation specific factor 5, 25 kD subunit	NM_018210	1.82
hypothetical protein FLJ10769	NM_006694	1.82
Jumping translocation breakpoint	AA634272	1.82
signal transducer and activator of transcription 3 (acute-phase response factor)	NM_005951	1.82
metallothionein 1H	BE964125	1.82
RNA isopentenylpyrophosphate transferase	NM_014178	1.82
HSPC158 protein	BC003689	1.82
high-mobility group (nonhistone chromosomal) protein 17	NM_007002	1.82
cell membrane glycoprotein, 110000M(r) (surface antigen)	AA918224	1.82
KIAA0052 gene product	AF109873	1.82
CREBBP/EP300 inhibitory protein 1	D31888	1.82
KIAA0071 protein	AI971281	1.82
ATP citrate lyase	AA054734	1.82
Cip1-interacting zinc finger protein	AW612311	1.82
v-erb avian sarcoma virus CT10 oncogene homolog	NM_024313	1.82
hypothetical protein MGC3731	NM_003683	1.82
DNA segment on chromosome 21 (unique) 2058 expressed sequence	BC000519	1.82
RuvB-like 1 (E. coli)	AL567295	1.82
adaptor-related protein complex 2, beta 1 subunit	AL031313	1.82
eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)	AI375486	1.81
adenomatosis polyposis coli	AB036737	1.81
arginine-glutamic acid dipeptide (RE) repeats	AF277194	1.81
putative membrane protein	X74070	1.81
basic transcription factor 3	AA320784	1.81
ribosomal protein S10	AL133228	1.81
Consensus includes gb:AL133228/DEF=Human DNA sequence from clone RPS-1071L10 on chromosome 20 Corians part of a gene for a new member of the thymosininterle	NM_002811	1.81
proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog)	NM_001662	1.81
ADP-ribosylation factor 5	L18964	1.81
protein kinase C, iota	NM_013245	1.81
vacuolar protein sorting factor 4A	NM_006759	1.81
UDP-glucose pyrophosphorylase 2	NM_004317	1.81
araX (bacterial) arsenite transporter, ATP-binding, homolog 1	NM_018351	1.81
hypothetical protein FLJ11183	NM_014944	1.81
KIAA0811 protein	NM_017864	1.81
hypothetical protein FLJ20530	NM_014939	1.80
KIAA1012 protein	AU154358	1.80
KIAA0217 protein	BC000830	1.80
staufen, RNA binding protein (Drosophila)	NM_002137	1.80
heterogeneous nuclear ribonucleoprotein A2/B1	NM_005605	1.80
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calciurexin A gamma)	U36945	1.80
cyclic-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)	NM_006353	1.80
high-mobility group (nonhistone chromosomal) protein 17-like 3	AB018302	1.80
KIAA0759 protein	AI475965	1.80
glutamy-prolyl-RNA synthetase	NM_001087	1.80
angle-associated, migratory cell protein	NM_004832	1.80
glutathione-S-transferase like; glutathione transferase omega	AL527334	1.80
hypothetical protein DKFZ056A00523	AA388062	1.80
hypothetical protein FLJ11583	U95622	1.80
midustin 1	NM_015999	1.80
CGI-45 protein		

eukaryotic translation termination factor 1	M75715	1.80
CpG binding protein	NM_014593	1.79
erbB2 interacting protein	NM_018895	1.79
hypothetical protein FLJ11198	NM_018358	1.79
putative receptor protein	NM_003878	1.79
DNA segment on chromosome X (unique) 9879 expressed sequence	NM_006014	1.79
RecD protein-like (DNA helicase Q1-like)	BC001052	1.79
DKFZP564D116 protein	BE862354	1.79
chromosome 8 open reading frame 11	BC000388	1.79
CGI-38 protein	NM_015963	1.79
nuclear receptor subfamily 1, group D, member 1	X72631	1.79
COP9 homolog	BC003090	1.79
inhibitor of DNA binding 3, dominant negative helix-loop-helix protein	NM_002167	1.79
CGI-31 protein	NM_015959	1.79
SET translocation (myeloid leukemia-associated)	A276616	1.79
leucine-rich PPR-motif containing	M92439	1.79
HLA class II region expressed gene KE2	NM_014280	1.79
myosin IE	NM_004998	1.79
chromosome 11 hypothetical protein ORF3	NM_020154	1.79
butyrophilin, subfamily 3, member A2	BC002832	1.79
ribosomal protein L17	NM_000985	1.79
splicing factor 3b, subunit 2, 145KD	NM_008842	1.79
KIAA0217 protein	AW628886	1.79
sorting nexin 1	NM_003099	1.79
phosphoglycerate dehydrogenase	NM_008623	1.78
MIL1 protein	W84482	1.78
polydioxin cofactor synthesis 2	NM_015367	1.78
isocitrate dehydrogenase 3 (NADP+) beta	NM_004531	1.78
putative protein similar to nussy (Drosophila)	NM_006899	1.78
protein kinase C, nu	NM_005768	1.78
short-chain dehydrogenase/reductase 1	NM_005813	1.78
ligase I, DNA, ATP-dependent	NM_004753	1.78
TF53 target gene 1	NM_000234	1.78
non-canonical ubiquitin conjugating enzyme 1	AB007458	1.78
uncharacterized bone marrow protein BMD42	AF151039	1.78
KIAA0522 protein	NM_016456	1.78
nuclear receptor subfamily 5, group A, member 2	BF571239	1.78
peroxisomal membrane protein 2 (22KD)	AF146343	1.78
hypothetical protein FLJ10287	NM_018663	1.78
aryl hydrocarbon receptor interacting protein	NM_018049	1.78
bromodomain-containing 4	NM_003977	1.78
MCM4, minichromosome maintenance deficient 4 (S. cerevisiae)	BF718610	1.78
hypothetical protein MGC10963	AA904621	1.78
breifeldin A-inhibited guanine nucleotide-exchange protein 1	NM_030573	1.77
hypothetical protein DKFZP564K0822	AK025637	1.77
uncharacterized hematopoietic stem/progenitor cells protein MDS032	NM_030796	1.77
single-stranded DNA-binding protein 1	NM_018467	1.77
UDP-Gal:beta-GlcNAc beta 1,4-galactosyltransferase, polypeptide 4	NM_003143	1.77
gp.NM_014128.1 /DEF=Homo sapiens PRO0470 protein (PRO0470), mRNA, /FEA=mRNA (GEN=PRO0470) PROD=PRO0470 protein /DB_XREF=gj:7662555 /UG=Ha.278931	BC004523	1.77
KIAA0573 protein	NM_014128	1.77
leucine-rich PPR-motif containing	BC006374	1.77
tubulin tyrosine ligase-like 1	A653608	1.77
heterogeneous nuclear ribonucleoprotein C (C1/C2)	NM_012263	1.77
nuclear transcription factor Y, gamma	AV725195	1.77
guanine nucleotide binding protein (G protein), beta polypeptide 1	BC005003	1.77
hypothetical protein	AF070603	1.77
complement component 1, q subcomponent binding protein	BF038366	1.77
hypothetical protein MGC5244,	AU151801	1.77
DnaJ (Hsp40) homolog, subfamily B, member 12	NM_031213	1.77
hypothetical protein	NM_017626	1.77
SCAN domain-containing 1	NM_015705	1.77
zinc finger protein 259, pseudogene	NM_016558	1.77
	Z95118	1.77

Interferon, alpha 21	M12350	1.77
aryl hydrocarbon receptor interacting protein	AL556532	1.77
Integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4 receptor)	NM_000885	1.77
arginine-rich, mutated in early stage tumors	NM_006010	1.77
tubulin-interacting protein	AL080147	1.77
ATPase, aminophospholipid transporter (APLT), Class 1, type 8A, member 1	AB013452	1.77
hypothetical protein FLJ22104	AK000684	1.77
lipase A, lysosomal acid, cholesterol esterase (Wolman disease)	NM_000235	1.77
pumilio homolog 1 (Drosophila)	BG474429	1.76
X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD)	AA205834	1.76
HCR (e-helix coiled-coil rod homologue)	AF588988	1.76
APCS autophagy 5-like (S. cerevisiae)	AF293841	1.76
hypothetical protein FLJ21908	NM_024604	1.76
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide	NM_003406	1.76
phosphoprotein regulated by mitogenic pathways	NM_025195	1.76
leucine-rich and death domain containing	AF274972	1.76
SH3-domain, GRB2-like, endophilin B2	AW131863	1.76
glutathione synthetase	L42531	1.76
cytoplasmic polysenylation element binding protein	NM_030594	1.76
mitochondrial ribosomal protein L42	BE782148	1.76
regulator of G-protein signalling 10	AI744627	1.76
cyclin G2	L49506	1.76
hypothetical protein DKFZ434G0522	NM_017566	1.76
res-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)	BG292367	1.76
phosphomethyltransferase kinase	NM_006556	1.76
small acidic protein	NM_014267	1.76
LAT1-3TM protein	NM_031211	1.76
BCS1-like (yeast)	NM_004328	1.76
DKFZP564J0123 protein	BC002873	1.75
sorbitol dehydrogenase	L29008	1.75
pituitary tumor-transforming 1 interacting protein	NM_004339	1.75
thymosin, beta 10	NM_021103	1.75
serine protease inhibitor, Kunitz type, 2	AF027205	1.75
hypothetical protein FLJ14040	AK024102	1.75
signal recognition particle 72kD	AF069765	1.75
homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease	NM_014285	1.75
transcriptional adaptor 3 (ADA3, yeast homolog)-like (PCAF histone acetylase complex)	AK024982	1.75
protein phosphatase 4 (formerly X), catalytic subunit	BC001416	1.75
ribosomal protein S3A	NM_001006	1.75
DKFZP586O0120 protein	NM_014077	1.75
H3 histone family, member H	NM_003534	1.75
clone FLB4739	AA133341	1.75
Consensus includes gb:AF064606.1 /DEF=Homo sapiens KB07 protein mRNA, partial cds. /FEA=mRNA /PROD=KB07 protein /DB=Hs.306242 Homo	NM_005065	1.75
sef-1 suppressor of lin-12-like (C. elegans)	AL060112	1.75
Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722), /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.33273	NM_018834	1.75
matrin 3	AV702810	1.75
SET translocation (myeloid leukemia-associated)	NM_025146	1.75
hypothetical protein FLJ13194	NM_025146	1.75
gb:Z25421.1 /DEF=H. sapiens protein-serine/threonine kinase gene, complete cds. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405720 /FL=gb.Z25421.1	NM_003993	1.75
CDC-like kinase 2	AV700224	1.75
casein kinase 1, delta	NM_001021	1.75
ribosomal protein S17	NM_004645	1.75
collin	U38980	1.75
postmeiotic segregation increased 2-like 11	NM_004699	1.75
DNA segment on chromosome X (unique) 9928 expressed sequence	BC002492	1.75
hypothetical protein FLJ20244	D83778	1.75
KIAA0194 protein	AB028628	1.75
DKFZP547E2110 protein	AC004544	1.75
cytochrome c oxidase subunit VIIa polypeptide 2 (liver)	BE881529	1.75
KIAA0731 protein	AF053970	1.75
outer dense fibre of sperm tails 2	NM_001489	1.75
GLE1 RNA export mediator-like (yeast)	NM_014019	1.75
HSPC009 protein		

numb homolog (Drosophila)	AF015040	1.75
vascular endothelial growth factor	AF022375	1.75
ribosomal protein L36a	NM_001001	1.75
protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I)	NM_000845	1.75
lactate dehydrogenase C	NM_002301	1.75
serine/threonine kinase 19	NM_004197	1.75
ES1 (zebrafish) protein, human homolog of	NM_004649	1.75
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3	J03198	1.74
hypothetical protein F23985	BF215644	1.74
nif endonuclease III-like 1 (E. coli)	NM_016404	1.74
NAD(P) dependent steroid dehydrogenase-like; H105e3	U79718	1.74
membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10)	BC000245	1.74
H2B histone family, member A	NM_007287	1.74
RING1 and YY1 binding protein	BC001131	1.74
nuclear transcription factor, X-box binding 1	AB028551	1.74
N-sulfoglucosamine sulfohydrolase (sulfamidase)	AW291398	1.74
adaptor-related protein complex 2, alpha 2 subunit	U30894	1.74
RPB5-mediated protein	AI25280	1.74
DKFZP434D1335 protein	AB006572	1.74
ribosomal protein L12	AL117499	1.74
pseudouridylyl synthase 1	NM_000976	1.74
phosphatidylinositol 4-kinase, catalytic, alpha polypeptide	NM_025215	1.74
G-2 and S-phase expressed 1	NM_002650	1.74
culin 4B	BF306380	1.74
methylenetetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase	AI650819	1.74
G protein-coupled receptor 68	NM_006638	1.73
paired-like homeodomain transcription factor 1	U35398	1.73
Consensus includes gb:AU144068 /FEA=EST /DB_XREF=est:AU144066 /CLONE=HEMBA1000798 /UG=Hs.285519 Homo sapiens cDNA FLJ11804	NM_002653	1.73
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1	AU144066	1.73
HSPC015 protein	NM_001055	1.73
tumor necrosis factor receptor superfamily, member 6b, decoy	NM_014026	1.73
muS homolog 3 (E. coli)	J04810	1.73
serine/threonine kinase 17b (apoptosis-inducing)	NM_004226	1.73
tyrosyl-RNA synthetase	AW245400	1.73
RAD50 homolog (S. cerevisiae)	U63139	1.73
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD	AI882255	1.73
myeloid cell leukemia sequence 1 (BCL2-related)	NM_021960	1.73
RNA binding motif protein 5	NM_005778	1.73
Consensus includes gb:BE544683 /FEA=EST /DB_XREF=gi:9773308 /DB_XREF=est:601074131F1 /CLONE=IMAGE:3460399 /UG=Hs.110407 ESTs, Weakly similar to coded	BC000120	1.73
general transcription factor IIF, polypeptide 1 (74kD subunit)	BF971587	1.73
tubulin, beta polypeptide	AI195589	1.73
hypothetical protein DKFZP434E1723	NM_012234	1.73
RING1 and YY1 binding protein	NM_002389	1.73
membrane cofactor protein (CD48, trophoblast-lymphocyte cross-reactive antigen)	AU135154	1.73
a disintegrin and metalloproteinase domain 10	O25304	1.73
Rac/Cdc42 guanine exchange factor (GEF) 6	NM_020123	1.73
SM-11044 binding protein	AA827892	1.72
Cluster incl. AA827892:cd08604.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:1387358 /clone_end=3' /gb=AA827892 /gi=2900255 /ug=Hs.239016 /len=622	NM_016062	1.72
CDi-128 protein	BC001604	1.72
hematopoietic protein 1	NM_001030	1.72
ribosomal protein S27 (metalloproteinin 1)	AB018272	1.72
KIAA0773 protein	NM_018131	1.72
hypothetical protein FLJ10540	NM_002583	1.72
PRKC, apoptosis, WT1, regulator	NM_004271	1.72
MD-1, RP105-associated	NM_006808	1.72
protein translocation complex beta	L23928	1.72
serine hydroxymethyltransferase 1 (soluble)	AA628948	1.72
adenylosuccinate synthase	U37012	1.72
cleavage and polyadenylation specific factor 1, 160kD subunit	NM_003720	1.72
Down syndrome critical region gene 2	S60876	1.72
zinc finger protein, subfamily 1A, 1 (Ikaros)	AW025150	1.72
mitogen-activated protein kinase kinase kinase 12		

putative nuclear protein	AF281030	1.72
CGI-146 protein	AF229834	1.72
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AK026573	1.72
putative neurotransmitter receptor	NM_003967	1.72
proteasome (prosome, macropain) subunit, beta type, 2	NM_002794	1.72
ribosomal protein L30	L05095	1.72
gb:Z25430.1 /DEF=H sapiens protein-serine/threonine kinase /DB_XREF=gi:405738 /FL=gb:Z25430. Z25430		1.72
MAP/microtubule affinity-regulating kinase 3	A1745639	1.71
methionine-RNA synthetase	AA621558	1.71
methylthioadenosine phosphorylase	AF109294	1.71
oxoglutarate (alpha-ketoglutarate) dehydrogenase (lipoamide)	NM_002541	1.71
small nuclear ribonucleoprotein D1 polypeptide (16kD)	BC001721	1.71
loricrin	NM_000427	1.71
EAP30 subunit of ELL complex	NM_007241	1.71
R0D1 regulator of differentiation 1 (S. pombe)	AW190873	1.71
L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain	AF001903	1.71
hypothetical protein hCLA-iso	NM_030837	1.71
likely ortholog of mouse tubulin alpha 4	NM_025019	1.71
transporin-SR	AF145029	1.71
Consensus includes gb:AK001816.1 /DEF=Homo sapiens cDNA FLJ10854 fts, clone PLACE1000383, highly similar to Homo sapiens mRNA for MTMR1 protein. /FEA=mRNA	AK001816	1.71
alcohol dehydrogenase 5 (class III), chi polypeptide	M29872	1.71
ATP citrate lyase	NM_001096	1.71
tumor suppressor deleted in oral cancer-related 1	NM_005851	1.71
chromosome 14 open reading frame 2	AF116639	1.71
protein tyrosine phosphatase, receptor type, N	NM_028246	1.71
hypothetical protein FLJ22301	NM_024836	1.71
interleukin 4 receptor	NM_000418	1.71
proteasome (prosome, macropain) activator subunit 1 (PA28 alpha)	NM_06283	1.71
calcium and integrin binding protein (DNA-dependent protein kinase interacting protein)	NM_06384	1.71
gb:NM_006328.1 /DEF=Homo sapiens seven transmembrane domain protein (NIFE14), mRNA. /FEA=mRNA GEN=NIFE14 /PROD=seven transmembrane domain protein /C	NM_006326	1.71
B-cell CLL/lymphoma 6 (zinc finger protein 51)	NM_001706	1.71
myosin, light polypeptide 4, alkali; atrial; embryonic	NM_001706	1.71
KIAA0785 protein	AF116676	1.71
NADH dehydrogenase (ubiquinone) Fe-S protein 5 (15kD) [NADH-coenzyme Q reductase]	AA913513	1.71
cytochrome c oxidase subunit VIa polypeptide 1	NM_004552	1.71
gb:AF116705.1 /DEF=Homo sapiens PRO2489 mRNA, complete cds. /FEA=mRNA /PROD=PRO2489 /DB_XREF=gi:7959908 /FL=gb:AF116705.1	NM_004373	1.71
M-phase phosphoprotein 9	AF116705	1.71
polymerase (DNA-directed) sigma	A1990326	1.71
ataxia telangiectasia mutated (includes complementation groups A, C and D)	NM_006899	1.71
small nuclear RNA activating complex, polypeptide 3, 50kD	NM_000051	1.70
aradine homolog 2 (Drosophila)	NM_006321	1.70
transformer-2 alpha (trh-2 alpha)	NM_078896	1.70
Cluster Incl. AA150503.z08g04.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-491768 /clone_end=3 /gb=AA150503 /gi=1722017 /ug=Hs.24435 /len=631	AA150503	1.70
Cluster Incl. AA150503.z08g04.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-491768 /clone_end=3 /gb=AA150503 /gi=1722017 /ug=Hs.24435 /len=631	J03620	1.70
dihydropyrimidine dehydrogenase (E3 component of pyruvate dehydrogenase complex, 2-oxo-glutarate complex, branched chain keto acid dehydrogenase complex)	A1885109	1.70
bromodomain-containing 7	BE966695	1.70
hypothetical protein FLJ14393	NM_005720	1.70
actin related protein 2/3 complex, subunit 1B (41 kD)	NM_007363	1.70
non-POU-domain-containing, octamer-binding	NM_004925	1.70
equopoin 3	NM_018281	1.70
STE20-like kinase	NM_020365	1.70
eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD)	NM_015870	1.70
gb:NM_015870.1 /DEF=Homo sapiens endogenous retrovirus H D1 leader region/integrase-derived ORF1, ORF2, and putative envelope protein (HSU88895), mRNA. /FEA=ml	A1139589	1.70
SWAP-70 protein	AA628140	1.70
Cluster Incl. AA628140.z07e06.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-1055746 /clone_end=3 /gb=AA628140 /gi=2540527 /ug=Hs.126089 /len=302	BG391005	1.70
Kalao	NM_015523	1.70
small fragment nuclease	AL050361	1.70
mitochondrial ribosomal protein S18B	NM_024040	1.70
hypothetical protein MGC2491	BC001312	1.70
protein disulfide isomerase-related protein	AK000970	1.70
similar to S. cerevisiae SSMA4	AK021780	1.70
phosphatidyserine receptor	AC005943	1.70
ubiquinol-cytochrome c reductase (6.4kD) subunit	AF005422	1.70
chloride channel, nucleotide-sensitive, 1A		

nuclease sensitive element binding protein 1	BC002411	1.69
general transcription factor (TF, polypeptide 1 (74KD subunit))	AW190445	1.69
hypothetical protein	NM_019005	1.69
P450 (cytochrome) oxidoreductase	AF25834.1	1.69
S100 calcium binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog)	NM_002961	1.69
ELKS protein	A8029004	1.69
hypothetical protein	NM_015932	1.69
hypothetical protein FLJ13984	NM_024770	1.69
ras homolog gene family, member G (rho G)	NM_001665	1.69
ribosomal protein L7	NM_000971	1.69
sialidase 1 (lysosomal sialidase)	U84246	1.69
KIAA0746 protein	A8018289	1.69
coatamer protein complex, subunit alpha	A1621079	1.69
hypothetical protein FLJ22940	NM_024571	1.69
transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C)	NM_005648	1.69
pM5 protein	N39536	1.69
sortilin-related receptor, L (OLR class) A repeats-containing	NM_003105	1.69
hypothetical protein MGC2668	NM_030914	1.68
ribosomal protein L21	NM_000982	1.68
RAN binding protein 3	NM_018446	1.68
glycosyltransferase AD-017	NM_016096	1.68
stearyl-CoA desaturase (delta-9-desaturase)	A8032261	1.68
hypothetical protein FLJ22795	AB69052	1.68
HSPCQ38 protein	NM_016096	1.68
ADP-ribosylation factor related protein 1	AF230401	1.68
hypothetical protein MGC10753	NM_003224	1.68
galactokinase 2	NM_016628	1.68
hypothetical protein FLJ11773	NM_002044	1.68
U2(RNU2) small nuclear RNA auxiliary factor 1 (non-standard symbol)	NM_021934	1.68
embryonic ectoderm development	NM_008758	1.68
hypothetical protein FLJ11896	AF080227	1.68
hypothetical protein MGC2731	NM_024611	1.68
mitochondrial carrier homolog 1	NM_024068	1.68
proteasome (prosome, macropain) subunit, beta type, 6	AF189289	1.68
cytochrome c oxidase subunit VIb pseudogene-3	BC000835	1.68
KIAA0999 protein	AL031594	1.67
Consensus includes gb:N95443 /FEA=EST /DB_XREF=gi:1287753 /DB_XREF=est:zb81c12.s1 /CLONE=IMAGE-310006 /UG=Hs.19180 Homo sapiens mRNA; cDNA DKFZp5-N95443	AA044154	1.67
B-cell receptor-associated protein BAP29	NM_018844	1.67
S-adenosylmethionine decarboxylase 1	M21154	1.67
KIAA0871 protein	AF112221	1.67
Consensus includes gb-AV708130 /FEA=EST /DB_XREF=gi:10725395 /DB_XREF=est:AV708130 /CLONE=ADCAMA05 /UG=Hs.283965 Homo sapiens scavenger receptor cti	AF143684	1.67
myosin IXB	BC005212	1.67
hypothetical protein FLJ20003	NM_013366	1.67
anaphase-promoting complex subunit 2	BC005161	1.67
hypothetical protein MGC4638	NM_005340	1.67
histidine triad nucleotide binding protein	NM_023005	1.67
bromodomain adjacent to zinc finger domain, 1B	M18003	1.67
ferredoxin 1	BC002585	1.67
RAB7, member RAS oncogene family-like 1	NM_025233	1.67
nucleotide binding protein	S72931	1.67
c8.1A	NM_014418	1.67
T-cell leukemia/lymphoma 6	NM_006489	1.67
NS1-binding protein	NM_014183	1.67
HSPC162 protein	AL553254	1.66
DEAD(H) (Asp-Glu-Ala-Asp/His) box polypeptide 19 (DBP5 homolog, yeast)	U47077	1.66
protein kinase, DNA-activated, catalytic polypeptide	NM_005507	1.66
cofilin 1 (non-muscle)	U56331	1.66
sarcoglycan, delta (35kD dystrophin-associated glycoprotein)	AF080157	1.66
conserved helix-loop-helix ubiquitous kinase	BC006479	1.66
hypothetical protein MGC2749	AB051480	1.66
hypothetical protein FLJ20719	A1186701	1.66
inhibitor of growth family, member 1-like		

nuclease sensitive element binding protein 1	X96666	1.66
ubiquitin fusion degradation 1-like	BC001049	1.66
ribosomal protein L13	BC004954	1.66
proline synthetase co-transcribed (bacterial homolog)	AL138816	1.66
G-rich RNA sequence binding factor 1	BF058465	1.66
ORF	NM_015853	1.66
uncoupling hydrogen decarboxylase	AW614435	1.66
19A24 protein	NM_021181	1.66
ribosomal protein L7	BG389744	1.66
hypothetical protein	AA463853	1.66
hyaluronan-mediated motility receptor (RHAMM)	U29343	1.66
chromatin assembly factor 1, subunit A (p150)	BF062223	1.66
hypothetical protein FLJ10975	NM_018288	1.66
KIAA0052 protein	D29841	1.66
spermine synthase	NM_004595	1.66
amplified in osteosarcoma	AI653900	1.66
ubiquitin specific protease 22	AA621731	1.66
hypothetical protein FLJ12587	AL136796	1.66
EH-domain containing 4	AF320070	1.66
platelet-activating factor acetylhydrolase, isoform 1b, alpha subunit (45kD)	L13387	1.66
hypothetical protein FLJ11021 similar to splicing factor, arginine/serine-rich 4	BE396879	1.66
ATP-binding cassette, sub-family C (CFTR/MRP), member 9	NM_005691	1.66
Consensus includes gb:AK026168.1 /DEF=Homo sapiens cDNA; FLJ22515 fls, clone HRC12122, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence. /FT AK026168	IF1 AK026168	1.66
SNARE protein	NM_006555	1.65
KIAA1118 protein	AB029041	1.65
karyopherin (importin) beta 3	AU148466	1.65
hypothetical protein FLJ22116	NM_024624	1.65
gene from NF2/meningioma region of 22q12	BC003815	1.65
suppression of tumorigenicity 7	NM_018412	1.65
ELK1, member of ETS oncogene family	M25269	1.65
RAP1B, member of RAS oncogene family	NM_015646	1.65
poly(ADP-ribose) polymerase 1	X62006	1.65
hypothetical protein FLJ11475	NM_003899	1.65
Consensus includes gb:AU144378 /FEA=EST /DB_XREF=gi:1005899 /DB_XREF=gi:1005899 DB_XREF=gi:1005899 /DB_XREF=gi:1005899	FLJ11475	1.65
cytosine-rich protein 1 (intestinal)	NM_001311	1.65
SEC14-like 1 (S. cerevisiae)	NM_003003	1.65
zinc finger protein 217	NM_006526	1.65
RAD17 homolog (S. pombe)	AF096533	1.65
transcriptional intermediary factor 1	AL338264	1.65
HSPC159 protein	NM_014181	1.65
mitochondrial ribosomal protein L35	NM_016622	1.65
CGI-125 protein	NM_016060	1.65
hypothetical protein FLJ20756	NM_017952	1.65
KIAA0247 gene product	NM_014734	1.65
protein kinase D2	AL050147	1.65
DAZ associated protein 1	NM_018959	1.65
RAB3 GTPase-ACTIVATING PROTEIN	AI040009	1.65
ATPase, H+ transporting lysosomal (vacuolar proton pump), member M	AF100741	1.65
leucine-rich repeat-containing 2	NM_024512	1.65
chondroitin sulfate proteoglycan 6 (barnatan)	AI373676	1.65
zinc finger protein 216	AW471220	1.65
nuclear distributor gene C (A. nidulans) homolog	AF241788	1.65
aldo-keto reductase family 7, member A3 (afatoxin aldehyde reductase)	NM_012067	1.65
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (lprn), alpha 1	U22815	1.65
ceroid-lipofuscinosis, neuronal 5	AV700514	1.65
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (lprn), alpha 1	BG033593	1.65
hypothetical protein R0083M05.2	NM_021167	1.64
elastase 3A, pancreatic (protease E)	BC005918	1.64
growth factor, augment of liver regeneration (ERV1 homolog, S. cerevisiae)	AF124604	1.64
protease, serine, 25	NM_013247	1.64
branched chain aminotransferase 2, mitochondrial	NM_001190	1.64
low density lipoprotein receptor defect C complementing	NM_007357	1.64
MAX protein	AA723514	1.64

purinergic receptor P2X, ligand-gated ion channel, 2	NM_016318	1.64
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 5 (RNA helicase, 58kD)	NM_004398	1.64
microfilament-associated protein 2	NM_017459	1.64
hypothetical protein MGC11256	NM_024324	1.64
Consensus includes gb:AW304232 /FEA=EST /DB_XREF=gi:6713921 /DB_XREF=ast:vx82g01.x1 /CLONE=IMAGE:2825040 /UG=Hs.181357 laminin receptor 1 (67kD, riboso	AW304232	1.64
DNA (cytosine-5-methyltransferase 1	NM_001379	1.64
hypothetical protein FLJ10083	NM_017992	1.64
histidine triad nucleotide binding protein	U27143	1.64
inhibitor of growth family, member 3	NM_019071	1.64
elastase 3A, pancreatic (protease E)	CO0308	1.64
transcription factor A, mitochondrial	NM_012251	1.64
eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)	BC002719	1.64
ring finger protein 13	AF070558	1.64
hypothetical protein MGC4614	NM_024294	1.64
somatostatin receptor 5	NM_001053	1.64
interleukin enhancer binding factor 3, 90kD	BC032366	1.64
eukaryotic translation elongation factor 1 epsilon 1	NM_004280	1.64
CGI-96 protein	AL022316	1.64
histidine triad nucleotide binding protein	M33197	1.64
JTV1 gene	N32864	1.64
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	A1828526	1.64
hypothetical protein DC42	BC004383	1.64
exosome component Rrp41	NM_030921	1.64
H2A histone family, member M	AL039469	1.63
transducer of ERBB2, 1	NM_003513	1.63
actin, gamma 1	AA675892	1.63
hypothetical protein FLJ10439	BE741683	1.63
Consensus includes gb:AL049250.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D113 (from clone DKFZp564D113) /FEA=mRNA /DB_XREF=gi:4499989 /UG=Hs.194637	NM_018093	1.63
regulator of G-protein signalling 14	AF037195	1.63
chromosome 8 open reading frame 12	NM_022755	1.63
ash2 (absent, small, or homeotic-like (Drosophila)	AB020982	1.63
hypothetical protein	BF038366	1.63
metastasis-associated 1-like 1	BE222618	1.63
hypothetical protein FLJ11021 similar to splicing factor, arginine/serine-rich 4	NM_023012	1.63
ribosomal protein L12	AK026491	1.63
hypothetical protein FLJ12547	NM_024992	1.63
hypothetical gene MGC19595	W67887	1.63
phospholipase C, gamma 1 (formerly subtype 14b)	AL022394	1.63
prohibitin	AL560017	1.63
Consensus includes gb:AK025862.1 /DEF=Homo sapiens cDNA: FLJ22209 fs, clone HRC01496 /FEA=mRNA /DB_XREF=gi:10438505 /UG=Hs.288528 Homo sapiens cDNA	AK025862	1.63
zinc finger protein 79 (pT7)	X65232	1.63
tumor necrosis factor, alpha-induced protein 2	A1882445	1.63
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	AA215654	1.63
zinc finger protein 42 (myeloid-specific retinoic acid-responsive)	NM_003422	1.63
huntingtin interacting protein-1-related	AB014555	1.63
hypothetical protein FLJ12671	NM_030980	1.63
eukaryotic translation initiation factor 4A, isoform 2	NM_001987	1.63
aldolase C, fructose-bisphosphate	NM_005165	1.63
JTV1 gene	NM_006303	1.63
differentially expressed in FDCP 6 homolog (mouse)	NM_022047	1.63
DKFZp564J157 protein	NM_018457	1.63
putative dimethyladenosine transferase	BC002841	1.63
protein kinase C substrate 80K-H	NM_002743	1.63
MAGUK protein p55T: Protein Associated with Lins 2	NM_016447	1.63
hypothetical protein	AL098741	1.63
hypothetical protein PP2447	BC399562	1.63
mitogen-activated protein kinase phosphatase x	NM_020185	1.62
dihydropyrimidine S-acetyltransferase (E2 component of pyruvate dehydrogenase complex)	BF978872	1.62
proline arginine-rich end leucine-rich repeat protein	U41344	1.62
KIAA1028 protein	AB028951	1.62
FK506 binding protein 4 (59kD)	AA894574	1.62
aridase homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)	NM_005744	1.62

gb-NM_013307.1 /DEF=Homo sapiens non-functional folate binding protein (HSAF000381). mRNA. /FEA=CDS /GEN=HSAF000381 /PROD=non-functional folate binding prot NM_013307
 Consensus includes gb:AL049761 /DEF=Human DNA sequence from clone RP5-863C7 on chromosome 20p12.3-13. Contains the CSNK2A1 gene for casein kinase 2 alpha 1 AL049761
 enoyl Coenzyme A hydratase, short chain, 1, mitochondrial
 adaptor-related protein complex 2, alpha 2 subunit
 KIAA0737 gene product
 hypothetical protein
 zinc finger protein 268
 lactate dehydrogenase A
 MAD, mothers against decapentaplegic homolog 4 (Drosophila)
 UDP glycosyltransferase 1 family, polypeptide A1
 mitotic spindle coiled-coil related protein
 CD22 antigen
 polymyxin B resistance autoantigen 1 (75kD)
 calcium binding protein Cab45 precursor
 DKFZP664G2022 protein
 EH-domain containing 1
 integrin cytoplasmic domain-associated protein 1
 DEAD(H) (Asp-Glu-Ala-Asp/His) box polypeptide 3
 Consensus includes gb:A092978 /FEA=EST /DB_XREF=gi:5755693 /DB_XREF=IMAGE:2508513 /UG=Hs.288908 WAS protein family, member 2 /FL A092978
 brefeldin A-inhibited guanine nucleotide-exchange protein 2
 ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast)
 tubulin, alpha 1 (testis specific)
 similar to MRJ gene for a member of the DNAJ protein family (H. sapiens)
 high-mobility group (nonhistone chromosomal) protein 17-like 3
 HSPC128 protein
 CDC45 cell division cycle 45-like (S. cerevisiae)
 hypothetical protein FLJ20038
 zinc finger protein 161 homolog (mouse)
 DKFZP434J154 protein
 DKFZP586D0623 protein
 pinin, desmosome associated protein
 phosphoribosyl pyrophosphatase synthetase-associated protein 2
 v-rel reticuloendotheliosis viral oncogene homolog B, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3 (avian)
 nuclear factor (erythroid-derived 2)-like 1
 tubulin-specific chaperone e
 ribosomal protein S24
 carnitine palmitoyltransferase I, muscle
 TEO protein
 origin recognition complex, subunit 3-like (yeast)
 eukaryotic translation initiation factor 1A
 DKFZP586G011 protein
 E51 (zebrafish) protein, human homolog of
 sperm associated antigen 4
 Consensus includes gb:BE733979 /FEA=EST /DB_XREF=gi:10147971 /DB_XREF=est:601571007 /CLONE=IMAGE:3925800 /UG=Hs.82202 ribosomal protein L17
 intercellular adhesion molecule 2
 suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein)
 macrophage myristoylated alanine-rich C kinase substrate
 ubiquitination factor E4B (UFD3 homolog, yeast)
 similar to S. cerevisiae Sec6p and R. norvegicus rsec6
 ubiquinol-cytochrome c reductase complex (7.2 kD)
 transcription factor A, mitochondrial
 Consensus includes gb:AC007204 /DEF=Homo sapiens chromosome 19, BAC 2732339 (CIT-B-320G13) /FEA=CDS /DB_XREF=gi:4559317 /UG=Hs.284229 Homo sapiens chr AC007204
 CCR4-NOT transcription complex, subunit 3
 Consensus includes gb:AJ224869 /DEF=Homo sapiens CXCR4 /FEA=mRNA /DB_XREF=gi:3059119 /UG=Hs.89414 chemokine (C-X-C motif) AJ224869
 cyclin-dependent kinase 4
 KIAA0542 gene product
 staufen, RNA binding protein (Drosophila)
 growth arrest and DNA-damage-inducible, beta
 N-acetyltransferase, homolog of S. cerevisiae ARO1
 Consensus includes gb:AL021395 /DEF=Human DNA sequence from clone RP1-269M15 on chromosome 20q12.1-13.12 Contains a gene similar to peptidylprolyl isomerase (cyt AL021395)
 MAD, mothers against decapentaplegic homolog 2 (Drosophila)
 WW domain binding protein 2
 U79458

KIAA1069 protein
formin-binding protein 3
chaperonin containing TCP1, subunit 8B (zeta 2)
C3H-type zinc finger protein; similar to D. melanogaster muscibind B protein
pro-oncogene receptor inducing membrane injury gene
serine/threonine kinase 38
squamous cell carcinoma antigen recognised by T cells
chromosome 21 open reading frame 4
polymerase (DNA directed), delta 1, catalytic subunit (125kD)
Consensus includes gb:BE588184 /FEA=EST /DB_XREF=gi:9811904 /DB_XREF=est:601341722F1 /CLONE=IMAGE:3683860 /UG=Hs.200400 Human DNA sequence from B
general transcription factor IIE, polypeptide 2 (beta subunit, 34kD)
ribosomal protein S4, X-linked
Friend leukemia virus integration 1
HSPC028 protein
synaptogyrin 2
cat eye syndrome chromosome region, candidate 5
Consensus includes gb:A1972416 /FEA=EST /DB_XREF=gi:5769332 /DB_XREF=est:wr39610.x1 /CLONE=IMAGE:2490067 /UG=Hs.24385 Human hbc647 mRNA sequence
APG12 autophagy 12-like (S. cerevisiae)
chromosome 11 hypothetical protein ORF4
hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit
mitogen-activated protein kinase kinase 7
hypothetical protein FLJ10631
Consensus includes gb:A1687738 /FEA=EST /DB_XREF=gi:4899032 /DB_XREF=est:tp93g08.x1 /CLONE=IMAGE:2206910 /UG=Hs.699 peptidylprolyl isomerase B (cyclophilin A)
ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)
activating transcription factor 4 (tax-responsive enhancer element B67)
H2A histone family, member 2
plasminogen activator, urokinase receptor
metallothionein 1H
putative membrane protein
chromosome 14 open reading frame 1
Consensus includes gb:BC003542.1 /DEF=Homo sapiens, clone IMAGE:3611719, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3611719) /DB_XREF=BC003542
PO94 (processing of precursor, S. cerevisiae) homolog
gb:M87268.1 /DEF=Human Igm VDJ-region mRNA, complete cds. /FEA=CDS /GEN=IGHM /PROD=igm /DB_XREF=gi:186197 /FL=gb:M87268.1
KIAA0399 protein
tubulin, alpha, ubiquitous
DKFZP434B168 protein
Sec61 gamma
LIN-7 protein 3
KIAA0293 protein
ornithine decarboxylase antizyme inhibitor
nucleic acid diphosphate linked moiety X-type motif 6
hypothetical protein FLJ12786
hypothetical protein
myosin regulatory light chain interacting protein
v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)
splicing factor, arginine/serine-rich 2
benzodiazepine receptor (peripheral)
ribosomal protein L13
hypothetical protein FLJ12492
protein similar to E.coli yhdg and R. capsulatus nifH3
natriuretic peptide precursor A
mitogen-activated protein kinase kinase 5
hypothetical protein FLJ10359
fumarate hydratase
dihydropyrimidine S-succinyltransferase (E2 component of 2-oxo-glutarate complex)
hypothetical protein FLJ10377
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4C
Cluster Ind. AJ363375:q5312.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2015951 /done_end=3 /gb=AJ363375 /gi=4114996 /ug=Hs.179801 /len=470
c-src tyrosine kinase
thyroid hormone receptor interactor 8
skatulin silent mating type information regulation 2 homolog 7 (S. cerevisiae)
c-Mpl binding protein
ribosomal protein S6

AK022810 1.61
AA675900 1.61
NM_006584 1.61
NM_005757 1.60
BG538627 1.60
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NM_005146 1.60
NM_006134 1.60
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A049392 1.59
A1363375 1.59
NM_004383 1.59
A1694023 1.59
NM_016538 1.59
A1743740 1.59
NM_001010 1.59

v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neurofiblioblastoma derived oncogene homolog (avian)
 C3HC4-type zinc finger protein
 translational inhibitor protein p14.5
 ubiquitin specific protease 12
 EZIG2 protein
 KIAA0240 protein
 eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)
 similar to Caenorhabditis elegans protein C42C1.9
 glutamine-fructose-6-phosphate transaminase 1
 v-Ha-ras Harvey rat sarcoma viral oncogene homolog
 KIAA0170 gene product
 coronin, actin binding protein, 1C
 eukaryotic translation initiation factor 4B
 hypothetical protein
 Golgi apparatus protein 1
 protein kinase, AMP-activated, gamma 1 non-catalytic subunit
 Treacher Collins-Franceschetti syndrome 1
 hypothetical protein FLJ20808
 CGI-51 protein
 zinc finger protein 262
 hypothetical protein MGC3062
 BRCA1-interacting protein 1
 chondroitin sulfate proteoglycan BEHAB/brevican
 guanidinoacetate N-methyltransferase
 coxsackie virus and adenovirus receptor
 polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)
 SMC2 structural maintenance of chromosomes 2-like 1 (yeast)
 adaptor-related protein complex 2, alpha 2 subunit
 phosphotyrosinotid transfer protein
 inositol(myo)-1(or 4)-monophosphatase 2
 hypothetical protein MGC5509
 solute carrier family 25 (mitochondrial carrier; citrate transporter), member 1
 hypothetical protein PRO2730
 phosphoinositide 3-kinase, platelet
 solute carrier family 21 (organic anion transporter), member 8
 nucleosome assembly protein 1-like 1
 deam-associated protein
 neighbor of COX4
 KIAA0409 protein
 KIAA0182 protein
 Consensus includes gb:AL161956.1 /DEF=Homo sapiens mRNA: cDNA DKFZp761A17121 (from clone DKFZp761A17121). /FEA=mRNA /DB_XREF=gi:7328008 /UG=Hs.306
 BCL2-like 1
 binder of Arf Two
 interferon, alpha 13
 KIAA0462 protein
 hepatoma-derived growth factor (high-mobility group protein 1-like)
 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
 eukaryotic translation initiation factor 2B, subunit 1 (alpha, 26kD)
 tetrapeptide repeat domain 1
 ribosomal protein L12
 zinc finger protein 36, C3H type-like 2
 Cbort10 protein
 KIAA0555 gene product
 eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)
 serpin-specific protease
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10 (42kD)
 kinesin-associated protein 3
 HSPC125 protein
 hematopoietic PBX-interacting protein
 exonuclease 1
 Consensus includes gb:BG332462 /FEA=EST /DB_XREF=gi:13138900 /DB_XREF=est:602432837F1 /CLONE=IMAGE-4550505 /UG=Hs.334.3 phosphoglycerate dehydrogenase
 hypothetical protein FLJ10099
 fatty-acid-Coenzyme A ligase, long-chain 1

X03363
 NM_024835
 N54448
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 AL568449
 BE645771
 NM_005343
 AB73553
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 AF360549
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 NM_004394
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 AA206181
 AL161956
 NM_001191
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 NM_006900
 AB007831
 NM_004494
 AA768906
 NM_001414
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 A356398
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 NM_014790
 U78525
 NM_021627
 NM_004544
 NM_014970
 NM_014165
 A935162
 NM_003886
 BG332462
 NM_017894
 NM_001995

hypothetical protein FLJ13491	AI431802	1.57
MAD1, midline arrest deficient-like 1 (yeast)	NM_003550	1.57
cytochrome c oxidase subunit VIc pseudogene 1	AF042165	1.57
cell division cycle 2-like 2	AL031282	1.57
hypothetical protein FLJ22490	BE044503	1.57
macrophage stimulating 1 (hepatocyte growth factor-like)	NM_020998	1.57
hydroxacyl glutathione hydrolase	NM_005326	1.57
KIAA0084 protein	D42043	1.57
peroxisomal short-chain alcohol dehydrogenase	NM_021004	1.57
CGI-18 protein	NM_015947	1.57
zinc finger protein 289, ID1 regulated	BE289871	1.57
thymopoleitin	BG391171	1.57
RAB7, member RAS oncogene family	BG281416	1.57
hypothetical protein MGC10763	AI560217	1.57
KIAA0943 protein	NM_013325	1.57
eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD)	AL031668	1.57
UDP-Gal4-betaGalNAc beta 1,4-galactosyltransferase, polypeptide 6	AF087159	1.57
replication factor C (activator 1) 1 (145kD)	L14922	1.57
methyl-CpG binding domain protein 4	AL558619	1.57
Consensus includes gb:AV733266 /FEA=EST /DB_XREF=gi:10850811 /DB_XREF=est:AV733266 /CLONE=cdAAJG04 /UG=Hs.76325 step II splicing factor SLU7	AV733266	1.57
Inositol polyphosphate-5-phosphatase, 145kD	NM_005541	1.57
CGG triplet repeat binding protein 1	NM_003663	1.57
RAD50 homolog (S. cerevisiae)	NM_005732	1.57
hypothetical protein MGC5468	AI589507	1.57
gb:AF293339.1 /DEF=Homo sapiens ornithine decarboxylase antizyme 4 mRNA, complete cds. /FEA=mRNA /PROD=ornithine decarboxylase antizyme 4 /DB_XREF=gi:11024; /CLONE=IMAGE:4448789	AF293339	1.57
fibroblast growth factor 4 (heparin secretory transforming protein 1, Kaposi sarcoma oncogene)	BF510715	1.57
nucleolin	NM_005381	1.57
isoprenylcysteine carboxyl methyltransferase	AL578502	1.57
low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)	BF304759	1.57
KIAA0650 protein	AA868754	1.57
hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor)	NM_001530	1.57
actin, beta	DS0604	1.57
splicing factor (CC1:3)	NM_004902	1.57
Consensus includes gb:G168283 /FEA=EST /DB_XREF=gi:12674986 /DB_XREF=est:602340822F /CLONE=IMAGE:4448789	BG168283	1.57
N-acetylglucosamine kinase	NM_017567	1.57
KIAA0251 protein	AK025504	1.57
Ewing sarcoma breakpoint region 1	AF327066	1.57
X-box binding protein 1	NM_005080	1.57
huntingtin interacting protein 2	NM_005339	1.57
U2 small nuclear ribonucleoprotein auxiliary factor (65kD)	NM_007279	1.57
Consensus includes gb:AL050217.1 /DEF=Homo sapiens mRNA; cDNA DKFZp58610523 (from clone DKFZp58610523). /FEA=mRNA /DB_XREF=gi:4884458 /UG=Hs.3084871	AL050217	1.57
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	AU138166	1.57
hypothetical protein KIAA1164	AL049226	1.57
zinc finger protein 22 (KIX 15)	AA744771	1.57
RAN binding protein 7	AA939270	1.56
hypothetical protein FLJ12910	NM_024573	1.56
ATPase, (Na+ /K+)-transporting, beta 4 polypeptide	AW975818	1.56
DKFZP586C0424 protein	NM_012069	1.56
cytochrome c oxidase subunit VIb pseudogene 1	AL050028	1.56
eIF4 associated protein HSPC021	AF042164	1.56
isocitrate dehydrogenase 3 (NAD+) gamma	NM_016091	1.56
glutaryl-Coenzyme A dehydrogenase	U69268	1.56
Rab geranylgeranyltransferase, beta subunit	NM_000159	1.56
hypothetical protein bK1048E9.5	U49245	1.56
KIAA0874 protein	NM_022081	1.56
protection of telomeres 1	AW572909	1.56
thioredoxin interacting protein	BC002923	1.56
hypothetical protein DKFZp564O0523	AA812232	1.56
lactate dehydrogenase B	AL136619	1.56
golgi associated, gamma adaptin ear containing, ARF binding protein 1	NM_002300	1.56
vesicle-associated membrane protein 3 (cellubrevin)	AW009695	1.56
serine/threonine kinase 12	NM_004781	1.56
	AB011446	1.56

putative GTP-binding protein similar to RAY/RAB1C	NM_006860	1.56
proteasome (prosome, macropain) 26S subunit, non-ATPase, 13	NM_002817	1.56
nuclear transcription factor Y, alpha	AL031778	1.56
amyloid beta (A4) precursor-like protein 2	BC004371	1.56
cyclin B2	NM_004701	1.56
cytochrome c oxidase subunit IV isoform 1	AA854966	1.56
ubiquitin specific protease 10	BC000263	1.56
uncharacterized hematopoietic stem/progenitor cells protein MDS026	A1138993	1.56
protein kinase, interferon-inducible double stranded RNA dependent activator	AF083033	1.56
BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIB (S.cerevisiae)	N91109	1.56
ADP-ribosylation factor 3	M33384	1.56
nuclear localization signal deleted in velocardiofacial syndrome	NM_003776	1.56
putative L-type neutral amino acid transporter	AW000954	1.56
polymyositis/scleroderma autoantigen 2 (100kD)	NM_002885	1.56
matrix metalloproteinase 11 (stromelysin 3)	NM_005940	1.56
DKFZP564K2062 protein	NM_015421	1.56
allantoinase	NM_018436	1.55
hypothetical protein FLJ12615 similar to membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 5)	NM_022474	1.55
protease, serine, 15	U02389	1.55
potassium voltage-gated channel, Shal-related subfamily, member 1	NM_004979	1.55
nascent-polypeptide-associated complex alpha polypeptide	NM_005594	1.55
keratin 6B	L42612	1.55
cytochrome c oxidase subunit Vb	AU57312	1.55
Mad4 homolog	AA831438	1.55
CDC5 cell division cycle 5-like (S. pombe)	AW268817	1.55
regulator of G-protein signalling 2, 24kD	NM_002923	1.55
hypothetical protein FLJ10808	NM_018227	1.55
protoporphyrinogen oxidase	NM_000309	1.55
transgelin 2	BC002616	1.55
regulator of G-protein signalling 10	NM_002925	1.55
KIAA0288 protein	AW205215	1.55
dabirin, light polypeptide (Lcb)	NM_001834	1.55
laminin, alpha 5	BC003355	1.55
translocase of inner mitochondrial membrane 17 homolog A (yeast)	BC004439	1.55
hypothetical protein MGC11279	NM_024326	1.55
LDL receptor adaptor protein	AA169780	1.55
tubulin, gamma 1	NM_001070	1.55
spermidine/spermine N1-acetyltransferase	BES71383	1.55
CtpX caseinolytic protease X homolog (E. coli)	NM_006660	1.55
hypothetical protein FLJ20568	AA438930	1.55
zinc finger protein 258	NM_007167	1.55
spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	U83867	1.55
hypothetical protein MGC20727	AU879381	1.55
macrophage stimulating 1 (hepatocyte growth factor-like)	U37055	1.55
cathepsin C	NM_001814	1.55
zinc finger protein 354A	NM_005649	1.55
palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile)	NM_000310	1.55
stearoyl-CoA desaturase (delta-9-desaturase)	AF116816	1.55
alpha integrin binding protein 63	NM_007216	1.55
hepatic leukemia factor	M95585	1.55
BAF53	NM_004301	1.55
cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)	NM_000079	1.55
A kinase (PRKA) anchor protein 13	M90360	1.55
tRNA-quinone transglycosylase	NM_031209	1.55
leucylcystinyl aminopeptidase	NM_005575	1.55
EH-domain containing 1	NM_006795	1.55
hypothetical protein	NM_016407	1.55
phosphatidylinositol binding clathrin assembly protein	AU722190	1.54
hypothetical protein MGC861	NM_024053	1.54
dCMP deaminase	AU856493	1.54
putative mitochondrial outer membrane protein import receptor	BG469030	1.54
inhibitor of DNA binding 2, dominant negative helix-loop-helix protein	D13891	1.54
MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	X63381	1.54

polycystic kidney disease 1 (autosomal dominant)	AK023376	1.54
capillary morphogenesis protein 1	AI610355	1.54
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1	NM_003079	1.54
uroporphyrinogen decarboxylase	M14016	1.54
pleckstrin homology-like domain, family A, member 1	AA576961	1.54
activating transcription factor 3	NM_001674	1.54
thioredoxin-like 2	AF118652	1.54
CD1A antigen, a polypeptide	M28825	1.54
PTD015 protein	NM_014040	1.54
NOL1R2 protein	AK021688	1.54
X-ray repair complementing defective repair in Chinese hamster cells 1	NM_006297	1.54
protein tyrosine phosphatase, receptor type, C	NM_002836	1.54
acyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase)	NM_001607	1.54
CGI-152 protein	NM_020410	1.54
nuclear factor (erythroid-derived 2)-like 2	NM_006164	1.54
GIM2 ganglioside activator protein	X62076	1.54
SMARIT1/DACT1 associated repressor protein	NM_015001	1.54
casein kinase 2, alpha 1 polypeptide	NM_001895	1.54
hypothetical protein FLJ12428	NM_022783	1.54
dyncin 4	AI038068	1.54
TCF3 (E2A) fusion partner (in childhood Leukemia)	NM_013342	1.54
Consensus includes gb:A742553/FEA=EST/DB_XREF=gi:5110841/DB_XREF=est:wg55d02.x1/CLONE=IMAGE:2368995/UG=Hs.184592 protein kinase, lysine deficient 1 / AI742553	AI742553	1.54
hypothetical protein FLJ10539	NM_018130	1.54
neuroblastoma RAS viral (v-ras) oncogene homolog	NM_002524	1.54
RNA binding motif protein 8A	AI738479	1.54
actin, gamma 1	BC026805	1.54
NADH dehydrogenase (ubiquinone) F ₀ -F ₁ protein 7 (20kD) (NADH-coenzyme Q reductase)	BC005954	1.54
formin-binding protein 17	AB011126	1.54
ubiquitously-expressed transcript	NM_004182	1.54
mitochondrial ribosomal protein L34	AB048652	1.53
copper chaperone for superoxide dismutase	NM_005125	1.53
superoxide dismutase 2, mitochondrial	W46388	1.53
homolog of yeast ribosome biogenesis regulatory protein RRS1	BC001811	1.53
ION3 protein	NM_015384	1.53
sialyltransferase 7D (alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase)	NM_014403	1.53
major histocompatibility complex, class I-like sequence	NM_001531	1.53
basigin (OK blood group)	AL550657	1.53
steroid dehydrogenase homolog	NM_016142	1.53
nuclear transcription factor Y, gamma	NM_014223	1.53
nucleolar protein GU2	AF334103	1.53
gb:BC002746.1/IDEF=Homo sapiens, Similar to dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase), clone MGC:3903, mRNA, complete cds. n	BC002746	1.53
hypothetical protein	AI081779	1.53
MAX protein	M64240	1.53
cyclin I	AF135162	1.53
ribosomal protein L14	AA638274	1.53
hypothetical protein FLJ20640	BC000890	1.53
programmed cell death 5	NM_004708	1.53
hypothetical protein FLJ10747	NM_018202	1.53
hypothetical protein FLJ20452	NM_017828	1.53
heterogeneous nuclear ribonucleoprotein K	BC000355	1.53
Consensus includes gb:AK022459.1/IDEF=Homo sapiens cDNA FLJ12397 fts, clone MAMMA1002769, weakly similar to Homo sapiens cell cycle progression restoration 8 prot	AK022459	1.53
PDGFA associated protein 1	NM_014481	1.53
protein kinase, lysine deficient 1	AI445745	1.53
EDF transcription factor 1	M96577	1.53
RNA binding protein	AF251062	1.53
metallothionein 1F (functional)	M10943	1.53
hypothetical protein DKF-Z6761H1710	NM_031297	1.53
Consensus includes gb:BE379008/FEA=EST/DB_XREF=gi:5324371/DB_XREF=est:801236272F1/CLONE=IMAGE:3608680/UG=Hs.99766 Homo sapiens mRNA; cDNA D	BE379008	1.53
malate dehydrogenase 2, NAD (mitochondrial)	BC001917	1.53
myogenic factor 3	BC000353	1.53
phosphatidylinositol 4-kinase, catalytic, beta polypeptide	NM_002851	1.53
CAMP responsive element binding protein 3 (luman)	AF028674	1.53
MCM3 mitochondrial maintenance deficient 3 (S. cerevisiae) associated protein	AJ010089	1.53

potassium channel, subfamily K, member 13	NM_022054	1.53
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	NM_003070	1.53
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	M60276	1.53
serine/threonine kinase 19	NM_003274	1.52
transmembrane protein 1	NM_000928	1.52
phospholipase A2, group IB (pancreas)	M57732	1.52
transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor	NM_000994	1.52
ribosomal protein L32	AF191653	1.52
nudix (nucleoside diphosphate linked moiety X)-type motif 4	NM_004896	1.52
vacuolar protein sorting 28 (yeast)	BC000966	1.52
gamma tubulin ring complex protein (76p gene)	NM_004922	1.52
SEC24 related gene family, member C (S. cerevisiae)	D84105	1.52
membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	NM_021971	1.52
GDP-mannose pyrophosphorylase B	NM_003377	1.52
vascular endothelial growth factor B	BC003375	1.52
mitochondrial ribosomal protein L3	U62136	1.52
ubiquitin-conjugating enzyme E2 variant 2	NM_001023	1.52
ribosomal protein S20	U67280	1.52
Consensus includes gb:AL109722.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIIMAGE 31619 /FEA=mRNA /DB_XREF=gi:5689814 /UG=Hs.170079 Homo sapiens	AF087853	1.52
calumenin	NM_014847	1.52
growth arrest and DNA-damage-inducible, beta	NM_001157	1.52
KIAA0144 gene product	BC004467	1.52
sinexin A11	NM_016821	1.52
KIAA0171 gene product	NM_002629	1.52
8-oxoguanine DNA glycosylase	NM_004890	1.52
phosphoglycerate mutase 1 (brain)	M36172	1.52
sperm associated antigen 7	NM_016172	1.52
myosin, light polypeptide 4, alkali; atrial, embryonic	NM_004941	1.52
putative glioblastoma cell differentiation-related	NM_022061	1.52
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 8 (RNA helicase)	T79953	1.52
Consensus includes gb:AK022061.1 /DEF=Homo sapiens cDNA FLJ11989 fls. clone HEMBB1001527 /FEA=mRNA /DB_XREF=gi:10433381 /UG=Hs.306622 Homo sapiens	D89646	1.52
KIAA0040 gene product	NM_008247	1.52
nucleoside homologue 6 (E. coli)	AF336851	1.52
protein phosphatase 3, catalytic subunit	NM_025230	1.52
hypothetical protein FLJ20596	NM_025230	1.52
hypothetical protein PRO2389	A4527238	1.52
nucleoporin 98KD	NM_006963	1.52
zinc finger protein 22 (KOX 15)	AF233437	1.52
myotubularin related protein 3	BC003080	1.52
peptidylprolyl isomerase F (cyclophilin F)	NM_000833	1.52
HSPC1171 protein	NM_014296	1.52
glutamate receptor, ionotropic, N-methyl D-aspartate 2A	NM_008661	1.52
calpain 7	AC004531	1.52
ribosomal protein L9	NM_006363	1.51
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 28	NM_002248	1.51
Sec23 homolog B (S. cerevisiae)	NM_021993	1.51
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1	NM_005332	1.51
FUS interacting protein (serine-arginine rich) 2	NM_001707	1.51
hemoglobin, zeta	NM_00940	1.51
B-cell CLL/lymphoma 78	AL562282	1.51
hypothetical protein MGC4276 similar to CG8198	D87453	1.51
hypothetical protein PP591	NM_022471	1.51
mitochondrial ribosomal protein S27	N36997	1.51
hypothetical protein FLJ13057 similar to germ cell-less	AK000145	1.51
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 16	NM_013239	1.51
Consensus includes gb:AK00145.1 /DEF=Homo sapiens cDNA FLJ20138 fls. clone COL07172 /FEA=mRNA /DB_XREF=gi:7020041 /UG=Hs.306384 Homo sapiens	M90355	1.51
protein phosphatase 2A 48 kDa regulatory subunit	AF006080	1.51
basic transcription factor 3, like 2	AU154985	1.51
growth hormone 2	BF026595	1.51
adenylyate kinase 2	AF172452	1.51
ribosomal protein S17	AJ236912	1.51
opioid growth factor receptor		
ADP-ribosyltransferase (NAD+-poly(ADP-ribose) polymerase)-like 2		

CGI-100 protein	NM_016040	1.51
Ins-related tyrosine kinase 3 ligand	NM_001459	1.51
hypothetical protein	NM_018988	1.51
esterase Diformylglutathione hydrolase	BC001169	1.51
hypothetical protein	NM_013298	1.51
phosphorylase kinase, beta	BG149218	1.51
cyclin B1	BE407516	1.51
NCK adaptor protein 1	NM_006153	1.51
spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	AL110273	1.51
hypothetical protein, clone Telent(hItaly_B41)_Strat02270_FL142	NM_020247	1.51
trichodeoid repeat containing 12	AL563727	1.51
X-ray repair complementing defective repair in Chinese hamster cells 4	AB017445	1.51
v-rel reticuloendotheliosis viral oncogene homolog (avian)	NM_002908	1.51
phospholipase A2-activating protein	AF145020	1.51
hypothetical protein FLJ20558	NM_017880	1.51
putative membrane protein	AB020980	1.51
chaperonin containing TCP1, subunit 2 (beta)	NM_006431	1.51
heterogeneous nuclear ribonucleoprotein A3	BG505670	1.51
hypothetical protein PRO1843	BF247371	1.51
CGI-48 protein	NM_016001	1.51
hypothetical protein MGC4054	NM_024341	1.51
CGI-07 protein	NM_015938	1.51
KIAA0415 gene product	AB007675	1.51
Consensus includes gb:W95043 /FEA=EST /DB_XREF=gi:1424182 /DB_XREF=est:zh46c11.s1 /CLONE=IMAGE:415124 /UG=Hs.79732 fibulin 1	W95043	1.51
FX506 binding protein 4 (S9KD)	NM_002014	1.51
G protein beta subunit-like	NM_022372	1.51
hypothetical protein FLJ22408	NM_024794	1.51
calreticulin	NM_004343	1.51
KIAA0478 gene product	NM_014856	1.51
hypothetical protein KIAA1164	AU121431	1.51
protein tyrosine phosphatase, non-receptor type 11	L07527	1.51
mitochondrial translational release factor 1	NM_004294	1.51
tumor protein 63 kDa with strong homology to p53	NM_003722	1.51
Wolf-Hirschhorn syndrome candidate 2	AF101434	1.51
YY1 transcription factor	Z14077	1.51
CCCTC-binding factor (zinc finger protein)	NM_006565	1.51
glutaminyl-peptide cyclotransferase (glutaminyl cyclase)	NM_012413	1.50
Consensus includes gb:AL515874 /FEA=EST /DB_XREF=gi:12779367 /DB_XREF=est:AL515874 /CLONE=CSODA001.YF06 (3 prime) /UG=Hs.83086 Homo sapiens GT212 m	AL515874	1.50
ubiquitin B	AB000635	1.50
Consensus includes gb:AL348009 /FEA=EST /DB_XREF=gi:4085215 /DB_XREF=est:qp61b09.x1 /CLONE=IMAGE:1927481 /UG=Hs.12421 hypothetical protein	NM_018955	1.50
cystathionase (cystathionine gamma-lyase)	AI348009	1.50
Consensus includes gb:BE871941 /FEA=EST /DB_XREF=gi:100322405 /DB_XREF=est:7a45g02.x1 /CLONE=IMAGE:3221810 /UG=Hs.157078 Homo sapiens cDNA FLJ127932	NM_001902	1.50
CGI-105 protein	BE871941	1.50
interleukin 2 receptor, gamma (severe combined immunodeficiency)	NM_016044	1.50
hypothetical protein FLJ22427	NM_000206	1.50
hypothetical protein FLJ21820	AK026080	1.50
peroxiredoxin 2	AK021672	1.50
pyrophosphatase (inorganic)	L19185	1.50
laminin, alpha 3 (nicotin (150KD), kalinin (165KD), epiligrin)	NM_021129	1.50
golgi associated, gamma adaptin ear containing, ARF binding protein 2	X77598	1.50
BCL2-associated athanogene 2	BE846414	1.50
16.7Kd protein	AF095192	1.50
hypothetical protein FLJ10154	NM_016139	1.50
hypothetical protein FLJ22570	NM_018011	1.50
Consensus includes gb:AL096828 /DEF=Human DNA sequence from clone RPS-963E22 on chromosome 20 Contains the 3 end of a novel gene similar to NY-REN-2 Antigen,	NM_024872	1.50
general transcription factor IIF, polypeptide 1 (74kD subunit)	AL096828	1.50
vaiA-RNA synthetase 2	NM_002096	1.50
hypothetical protein FLJ21613 similar to rat corneal wound healing related protein	BE780854	1.50
adaptor-related protein complex 1, beta 1 subunit	NM_021929	1.50
nucleosome assembly protein 1-like 1	NM_001127	1.50
hypothetical protein FLJ23320	AW148801	1.50
X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound	NM_024872	1.50
	AL023653	1.50

ribosomal protein S27a	NM_002954	1.50
dynamitin 1-like	NM_012062	1.50
HGF activator	NM_001528	1.50
small acidic protein	BE748698	1.50
KIAA0193 gene product	NM_014766	1.50
mitochondrial ribosomal protein L46	NM_022163	1.50
hypothetical protein FLJ10547	AL048795	1.50
colony stimulating factor 1 (macrophage)	M37435	1.50
nuclear receptor coactivator 3	U80737	1.50
DKFZP586J0119 protein	AJ011307	1.50
breast cancer 2, early onset	NM_000059	1.49
integrin, alpha X (antigen CD11C (p150), alpha polypeptide)	M81695	1.49
melastatin associated 1	BC006177	1.49
gag1 associated, gamma adaptin ear containing, ARF binding protein 1	NM_013365	1.49
chromosome 5 open reading frame 6	NM_016605	1.49
ATP-binding cassette, sub-family B (MDR/TPAP), member 8	NM_007188	1.49
E74-like factor 1 (ets domain transcription factor)	M82882	1.49
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polysaccharide-N-acetylglucosaminyl transferase)	NM_003605	1.49
hypothetical protein FLJ13114	H27225	1.49
M27830 Human 28S ribosomal RNA gene, complete cds (5', 3' represent transcript regions 5 prime, Middle, and 3 prime respectively)	M27830	1.49
KIAA0801 gene product	NM_014829	1.49
immediate early response 5	NM_016545	1.49
sorting nexin 17	NM_014748	1.49
capping protein (actin filament) muscle Z-line, alpha 2	AV65920	1.49
partner of RAC1 (arfpalin 2)	NM_012402	1.49
scratch homolog 1, zinc finger protein (Drosophila)	NM_031309	1.49
mitochondrial ribosomal protein S28	NM_014018	1.49
baculoviral IAP repeat-containing 5 (survivin)	NM_001188	1.49
H1 histone family, member 3	NM_005320	1.49
nucleolar protein (KKE/D repeat)	BE796327	1.49
L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain	NM_005327	1.49
aminolevulinatase, delta-, synthase 1	NM_006688	1.49
RAS p21 protein activator (GTPase activating protein) 1	M23812	1.49
hypothetical protein FLJ10307	NM_018053	1.49
2,4-dienoyl CoA reductase 1, mitochondrial	NM_001359	1.49
chromobox homolog 3 (HP1 gamma homolog, Drosophila)	BE748755	1.49
tumor protein p53-binding protein	AI150117	1.49
anaphase-promoting complex 1: meiotic checkpoint regulator	NM_022682	1.49
hypothetical protein	RS1077	1.49
hypothetical protein	NM_017571	1.49
H1 histone family, member 0	BC000145	1.49
hypothetical protein FLJ10450	NM_018095	1.49
retinoid acid receptor, alpha	T91506	1.49
sialyltransferase 7D ((alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase)	AB035172	1.49
tumor necrosis factor (ligand) superfamily, member 8	NM_001244	1.49
Consensus includes gb:AV681533 /FEA=EST /DB_XREF=gi:9882547 /DB_XREF=estAV681533 /CLONE=GLCGTD12 /UG=Hs.293637 ESTs	AV681533	1.49
dysactin 4 (p82)	NM_016221	1.49
Consensus includes gb:BE547674 /FEA=EST /DB_XREF=gi:9776319 /DB_XREF=est1601076255F1 /CLONE=IMAGE:3462325 /UG=Hs.204169 ESTs	BE547674	1.49
TH1-like (Drosophila)	NM_016397	1.49
sperm associated antigen 11	AF168619	1.49
nuclear factor, interleukin 3 regulated	NM_005384	1.49
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-W.	AB030077	1.49
hypothetical protein FLJ22637	NM_025165	1.49
progesterone receptor membrane component 1	AL547946	1.49
topoisomerase (DNA) III beta	D87012	1.49
serologically defined colon cancer antigen 8	U79271	1.48
serine/arginine repetitive matrix 2	X92110	1.48
HCGVIII-1 protein	NM_016333	1.48
endocytic receptor (macrophage mannose receptor family)	AB014609	1.48
Ran GTPase activating protein 1	NM_002883	1.48
prosaposin (variant Gaucher disease and variant melachromalic leukodystrophy)	NM_002776	1.48
programmed cell death 10	BC002506	1.48
RAN binding protein 16	H38643	1.48

putative mitochondrial outer membrane protein import receptor
golgin-67
three prime repair exonuclease 1
protein phosphatase 2 (formerly 2A), regulatory subunit B* (PR 72), alpha isoform and (PR 130), beta isoform
gb:AF081735.1 /DEF=Homo sapiens clone 023e08 My032 protein mRNA, complete cds, /FEA=mRNA /PROD=My032
BRCA2 and CDKN1A interacting protein
hypothetical protein FLJ20477
PEST-containing nuclear protein
pleckstrin homology, Sec7 and coiled/coil domains 2 (cytohesin-2)
zinc finger RNA binding protein
delta-like 3 (Drosophila)
sialoadhesin
H3 histone family, member F
Consensus includes gb:AK023938.1 /DEF=Homo sapiens cDNA FLJ13676 fts, clone THYRO1001401, /FEA=mRNA /DB_XREF=gi:10436033 /UG=Hs.287604 Homo sapiens c
SF115776
AL523310
AC002310
NM_016360
NM_004260
U55777
AF242521
NM_001920
A4479495
AK021557
NM_004539
AF072242
U33833
AF070655
NM_019082
A122099
A1220627
NM_024853
NM_003516
BC006379
AB019484
AC007856
NM_018054
AL137403
NM_000720
Consensus includes gb:AL137403.1 /DEF=Homo sapiens mRNA: cDNA DKF Zp434L092, /FEA=mRNA /DB_XREF=gi:6807951 /UG=Hs.306457 H AL137403
calcium channel, voltage-dependent, L type, alpha 1D subunit
Consensus includes gb:BF115739 /FEA=EST /DB_XREF=gi:10985215 /DB_XREF=est:7n64008.x1 /CLONE=IMAGE:3569246 /UG=Hs.15740 Homo sapiens mRNA: cDNA DK BF115739
heterogeneous nuclear ribonucleoprotein A3
Consensus includes gb:AW139502 /FEA=EST /DB_XREF=gi:5143220 /DB_XREF=est:UH-H-B11-aeq-e-08-Q-UI.s1 /CLONE=IMAGE:2720342 /UG=Hs.173108 Homo sapiens cDNA
hypothetical protein FLJ21016
Consensus includes gb:AL031602 /DEF=Human DNA sequence from clone RPS-1174N9 on chromosome 1p34.1-35.3. Contains the gene for a novel protein with IBR domain,
fizzled homolog 3 (Drosophila)
centromere protein A (17kD)
golgi SNAP receptor complex member 1
GTP cyclohydrolase 1 (dopa-responsive dystonia)
proteasome (prosome, macropain) 26S subunit, non-ATPase, 1
secretory carrier membrane protein 2
zinc finger protein 216
secreted protein of unknown function
cytochrome c oxidase subunit VIIa polypeptide 2 like
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)
DKFZp434J1813 protein
WD repeat domain 1
protein regulator of cytokinesis 1
signal recognition particle 72kD
interferon-related developmental regulator 2
Wilms' tumour 1-associated protein
hypothetical protein 394D8_6
BRCA1 associated RING domain 1
myosin, heavy polypeptide 9, non-muscle

VAMP (vesicle-associated membrane protein)-associated protein A (33kD)	AF154847	1.47
gene amplified in squamous cell carcinoma 1	AB037901	1.47
major histocompatibility complex, class I, F	A1689379	1.47
pre-mRNA splicing factor 17	NM_015991	1.47
Consensus includes gb:AF131823.1 /DEF=Homo sapiens clone 24997 mRNA sequence, /FEA=mRNA /DB_XREF=gi:4408659 /UG=Hs.308951 Homo sapiens clone 24997 mF	AF131823	1.47
Consensus includes gb:AF131823.1 /DEF=Homo sapiens clone 24997 mRNA sequence, /FEA=mRNA /DB_XREF=gi:4408659 /UG=Hs.308951 Homo sapiens clone 24997 mF	NM_012222	1.47
murY homolog (E. coli)	NM_021805	1.47
single Ig IL-1R-related molecule	AL138831	1.47
thioredoxin-like 2	AL031669	1.47
Consensus includes gb:AL031669 /DEF=Human DNA sequence from clone RPA-68120 on chromosome 20q11.23-12 Contains a gene for a protein similar to RPL12 (ribosomal protein L12)	NM_007097	1.47
catenin, light polypeptide (Lcb)	AW452623	1.47
Consensus includes gb:AW452623 /FEA=EST /DB_XREF=gi:5993399 /DB_XREF=est UI-H-B13-alu-d-09-0-UI.s1 /CLONE=IMAGE:3086609 /UG=Hs.14846 Homo sapiens mR	BC008481	1.47
tubulin, alpha, ubiquitous	L40027	1.47
glycogen synthase kinase 3 alpha	NM_003025	1.47
SH3-domain GRB2-like 1	NM_024810	1.47
hypothetical protein FLJ23018	A118402	1.47
hypothetical protein FLJ10539	AL117466	1.47
LIM domain kinase 2	NM_002131	1.47
high-mobility group (nonhistone chromosomal) protein isoforms 1 and Y	NM_004921	1.47
chloride channel, calcium activated, family member 3	NM_016013	1.47
COH45 protein	BF341845	1.47
casein kinase 1, alpha 1	AF227968	1.47
SH2-B homolog	Z97056	1.47
potassium inwardly-rectifying channel, subfamily J, member 4	AF193844	1.47
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 7A	NM_004819	1.47
sympkin; Huntingtin interacting protein 1	NM_004147	1.47
developmentally regulated GTP binding protein 1	NM_006626	1.47
zinc finger protein with interaction domain	NM_003588	1.47
culin 4B	BC001359	1.47
tyrosine 3-monooxygenase/tyrosinase 5-monooxygenase activation protein, beta polypeptide	NC01251	1.47
Consensus includes gb:AL344141 /FEA=EST /DB_XREF=gi:4081347 /DB_XREF=est tc02009.x1 /CLONE=IMAGE:2062673 /UG=Hs.229850 ESTs. Highly similar to G01251 R	AL344141	1.47
protein disulfide isomerase-related protein	NM_003742	1.47
adaptor-related protein complex 3, beta 1 subunit	NM_003664	1.47
zinc metalloproteinase (STE24 homolog, yeast)	NM_005857	1.47
hexokinase 1	NM_000188	1.47
B06 antigen	NM_016579	1.47
2-5'-oligoadenylate synthetase 2 (69-71 kD)	NM_016817	1.47
sorting nexin 7	NM_015976	1.47
cell division cycle 2-like 2	D13640	1.47
hypothetical protein FLJ20445	AL031282	1.47
Ras homolog enriched in brain 2	NM_017824	1.47
thyroid hormone receptor interactor 12	NM_005614	1.47
Era G-protein-like 1 (E. coli)	NM_004238	1.47
purinergic receptor P2X, ligand-gated ion channel, 7	AL562733	1.47
TNF receptor-associated factor 4	NM_002562	1.47
KIAA0678 protein	NM_004295	1.47
Cluster ind. AL050297; Homo sapiens mRNA; cDNA DKFp564N123 (from clone DKFp564N123) /cds=UNKNOWN /gb=AL050297 /lg=4886448 /ug=Hs.25894 /len=1340	AB014578	1.47
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5	AL050297	1.47
KIAA0332 protein	NM_006837	1.47
sorting nexin 3	A1184562	1.47
vascular endothelial growth factor	AB047360	1.46
zinc finger protein 131 (clone pTZ-10)	H95344	1.46
neighbor of A-kinase anchoring protein 95	AY966301	1.46
G protein-coupled receptor 87	NM_014371	1.46
nucleoporin 155KD	NM_023915	1.46
secretory carrier membrane protein 1	NM_004298	1.46
ADP37 protein	AF005037	1.46
ADP-ribosylation factor-like 1	AF260335	1.46
putative T1/ST2 receptor binding protein	BE890745	1.46
KIAA0246 protein	NM_006858	1.46
U5 snRNP-specific 40 kDa protein (hPrp8-binding)	NM_015136	1.46
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta	AL157420	1.46
heat shock 10kD protein 1 (chaperonin 10)	A1684894	1.46
	NM_002157	1.46

hypothetical protein 394D8_6
excision repair cross-complementing rodent repair deficiency complementation group 1 (includes overlapping antisense sequence)
aldol-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)
serine hydroxymethyltransferase 2 (mitochondrial)
putative nucleic acid binding protein RY-1
RAS protein activator like 1 (GAP1 like)
phosphorylase, glycogen; brain
ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)
Ran GTPase activating protein 1
amino-terminal enhancer of split
calpain, small subunit 1
gb:NM_005703.2 DEF=Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA, /FEA=CDS /GEN=UREB1 /PROD=upstream regulatory element bindin NM_005703
gb:AF092132.1 DEF=Homo sapiens PAK2 mRNA, complete cds, /FEA=mRNA /PROD=PAK2 /DB_XREF=gi:5138913 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete ABO40875
solute carrier family 7, (cationic amino acid transporter, y+ system) member 11
hypothetical protein
integrin-linked kinase-associated serine/threonine phosphatase 2C
phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminimidazole synthetase
ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-like 2
v-rat murine sarcoma 3811 viral oncogene homolog 1
SH3-domain, GRB2-like, endophilin B1
GRB2-associated binding protein 1
F-box only protein 9
ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)
RNA processing factor 1
proteasome (prosome, macropain) 26S subunit, ATPase, 5
transcription factor 2, hepatic; LF-83; variant hepatic nuclear factor
microtubule-associated protein 4
CD2 antigen (cytoplasmic tail) binding protein 2
antigen identified by monoclonal antibody Ki-67
hypothetical protein FLJ10241
casein kinase 2, alpha 1 polypeptide
HCR (e-helix coiled-coil rod homologue)
hypothetical protein FLJ23591
ATP-binding cassette, subfamily B (MDR/TAP), member 7
hypothetical protein FLJ12529
breast cancer 1, early onset
HIV-1 Tat interactive protein 2, 30 kDa
high density lipoprotein binding protein (vigilin)
hypothetical protein MGC14258
opoid growth factor receptor
chemokine-like factor 1
stromal antigen 2
actin A receptor, type IIB
hypothetical protein FLJ12443
glycogen synthase kinase 3 alpha
CCR4-NOT transcription complex, subunit 3
myozenin
cyclic AMP phosphoprotein, 19 kD
synapsin II
nucleosome assembly protein 1-like 1
zinc finger protein 91 (HPP7, HTF10)
arsenate resistance protein ARS2
ornithine decarboxylase antizyme 1
hypothetical protein FLJ10842
nuclear respiratory factor 1
Consensus includes gb:AK000345.1 /DEF=Homo sapiens cDNA FLJ20338 f1s, clone HEP12179, /FEA=mRNA /DB_XREF=gi:7020368 /UG=Hs.152677 Homo sapiens cDNA FI AK000345
PAL-1 mRNA-binding protein
putative c-Myc-responsive
hypothetical protein DKFZp566f2423
Consensus includes gb:AK026008.1 /DEF=Homo sapiens cDNA: FLJ22355 f1s, clone HRC06344, /FEA=mRNA /DB_XREF=gi:10438707 /UG=Hs.288283 Homo sapiens cDNA AK026008
v-myc myelocytomatosis viral oncogene homolog (avian)
synixin 7
casein kinase 1, gamma 3

BC001288 1.46
NM_001983 1.46
NM_003689 1.46
AW190316 1.46
BG252325 1.46
NM_004658 1.46
NM_002862 1.46
AA602532 1.46
BE378408 1.46
NM_001130 1.46
NM_001749 1.46
gb:NM_005703.2 DEF=Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA, /FEA=CDS /GEN=UREB1 /PROD=upstream regulatory element bindin NM_005703
gb:AF092132.1 DEF=Homo sapiens PAK2 mRNA, complete cds, /FEA=mRNA /PROD=PAK2 /DB_XREF=gi:5138913 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete ABO40875
NM_016458 1.46
AY024365 1.46
BE966876 1.46
NM_005484 1.46
NM_001654 1.46
AF257318 1.46
NM_002039 1.46
AI954041 1.46
NM_003348 1.46
NM_025065 1.46
AF055309 1.46
NM_000458 1.46
NM_002375 1.46
NM_006110 1.46
AU132185 1.46
AF768100 1.46
AI631874 1.46
AB029343 1.46
NM_025076 1.46
AB005289 1.46
NM_024811 1.46
AF005068 1.46
NM_006410 1.46
NM_005336 1.45
U79256 1.45
NM_007346 1.45
NM_016326 1.45
NM_008603 1.45
NM_001106 1.45
NM_024830 1.45
NM_019684 1.45
AF180474 1.45
NM_021245 1.45
BC003418 1.45
AW139618 1.45
AI888672 1.45
NM_003430 1.45
NM_015908 1.45
D87914 1.45
AJ278150 1.45
NM_005011 1.45
Consensus includes gb:AK000345.1 /DEF=Homo sapiens cDNA FLJ20338 f1s, clone HEP12179, /FEA=mRNA /DB_XREF=gi:7020368 /UG=Hs.152677 Homo sapiens cDNA FI AK000345
AF151813 1.45
AF040105 1.45
BE975139 1.45
AK026008 1.45
NM_002467 1.45
NM_003569 1.45
NM_004384 1.45

huntingtin interacting protein 1 1.45
 hypothetical protein FLJ12700 1.45
 succinate-CoA ligase, GDP-forming, beta subunit 1.45
 ADP-ribosylation factor-like 2 1.45
 replication protein A2 (32kD) 1.45
 phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase 1.45
 platelet-activating factor acetylhydrolase, isoform 1b, alpha subunit (45kD) 1.45
 hypothetical protein MGC2722 1.45
 cachectin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila) 1.45
 KIAA0637 gene product 1.45
 Consensus includes gb:AW371924 /FEA=EST /DB_XREF=gi:6876578 /DB_XREF=est:RC4-BT0311-251199-012-a07 /UG=Hs.313848 ESTs, Moderately similar to MGN_HUM_AW371924 1.45
 CGI-99 protein 1.45
 poly(pyrimidine) tract binding protein (heterogeneous nuclear ribonucleoprotein 1) 1.45
 bedlin 1 (coiled-coil, myosin-like BCL2 interacting protein) 1.45
 endoplasmic reticulum chaperone SIL1, homolog of yeast 1.44
 homolog of mouse quaking QKI (KH domain RNA binding protein) 1.44
 propionyl Coenzyme A carboxylase, beta polypeptide 1.44
 FUS interacting protein (serine-arginine rich) 1 1.44
 hypothetical protein FLJ21952 1.44
 polymerase (RNA) II (DNA directed) polypeptide E (25kD) 1.44
 interleukin 10 receptor, beta 1.44
 COP9 subunit 8 (MOV34 homolog, 34 kD) 1.44
 protein tyrosine phosphatase, non-receptor type 2 1.44
 Cluster Ind. AA160496:73a03.a1 Homo sapiens cDNA, 3 end /clone=IMAGE:592492 /clone_end=3 /gb=AA160498 /gi=1735963 /ug=Hs.110407 /len=542 1.44
 mitofusin 1 1.44
 splicing factor, arginine/serine-rich 5 1.44
 defensin, beta 2 1.44
 GM2 ganglioside activator protein 1.44
 general transcription factor IIIC, polypeptide 5 (63kD) 1.44
 VRK3 for vaccinia related kinase 3 1.44
 Cluster Ind. A1144299:qb59n06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:1704443 /clone_end=3 /gb=A1144299 /gi=3666108 /ug=Hs.106843 /len=765 1.44
 casein kinase 1, alpha 1 1.44
 Kruppel-like factor 12 1.44
 DKFZP434D156 protein 1.44
 phosphatidylinositol transfer protein, beta 1.44
 thiodoxin, mitochondrial 1.44
 host cell factor C1 (VP16-accessory protein) 1.44
 casein, kappa 1.44
 rat guanine nucleotide dissociation stimulator 1.44
 gb:NM_015930.1 /DEF=Homo sapiens vanilloid receptor-like protein 1 (VRL-1), mRNA. /FEA=mRNA /GEN=VRL-1 /PROD=vanilloid receptor-like protein 1 /DB_XREF=gi:77067 /NM_015930 1.44
 scaffold attachment factor B 1.44
 myelin protein zero-like 1 1.44
 hypothetical protein FLJ10546 1.44
 phosphatidylinositol 4-kinase, catalytic, alpha polypeptide 1.44
 Ndc68 (D. melanogaster)-like protein 1.44
 KIAA1253 protein 1.44
 regulator of nonsense transcripts 1 1.44
 Consensus includes gb:AL133053.1 /DEF=Homo sapiens mRNA; cDNA DKFZP434H1215 (from clone DKFZP434H1215); partial cds. /FEA=mRNA /GEN=DKFZP434H1215 /P AL133053 1.44
 Consensus includes gb:BF056534 /FEA=EST /DB_XREF=gi:10810430 /DB_XREF=est:7406609.x1 /CLONE=IMAGE:34413441 /UG=Hs.123469 ESTs, Weekly similar to AF208 /BF056534 1.44
 BCL2adenovirus E1B 19kD interacting protein 1 1.44
 Consensus includes gb:BC680028 /FEA=EST /DB_XREF=gi:10329604 /DB_XREF=est:60149326 /F1 /CLONE=IMAGE:38956407 /UG=Hs.84775 Human transposon-like element BEB08028 1.44
 Consensus includes gb:AW081820 /FEA=EST /DB_XREF=gi:6036972 /DB_XREF=est:xb55d12.x1 /CLONE=IMAGE:2580215 /UG=Hs.287317 ESTs, Weekly similar to AS9252 /AW081820 1.44
 poly(A) binding protein, cytoplasmic 4 (inducible form) 1.44
 polymerase (DNA directed), epsilon 1.44
 survival of motor neuron protein interacting protein 1 1.44
 DKFZp556F1019 protein 1.44
 golgi associated, gamma adaptin ear containing, ARF binding protein 2 1.44
 major histocompatibility complex, class II, DM beta 1.44
 capping protein (actin filament) muscle Z-line, beta 1.44
 Consensus includes gb:AU145293 /FEA=EST /DB_XREF=gi:11006814 /DB_XREF=est:AU145293 /CLONE=HEMBA1004405 /UG=Hs.61773 Homo sapiens cDNA FLJ11648 fl AU145293 1.44
 vesicle-associated membrane protein 2 (synaptobrevin 2) 1.44
 gb:NM_020382.1 /DEF=Homo sapiens PRSET domain containing protein 07 (SET07), mRNA. /FEA=mRNA /GEN=SET07 /PROD=PRSET domain containing protein 07 /DB_X_NM_020382 1.44

KIAA0540 protein	AB011112	1.44
ribosomal protein S23	NM_001025	1.44
KIAA0816 protein	AB014516	1.44
thionexin	AF313911	1.44
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	BC000409	1.44
signal transducer and activator of transcription 3 (acute-phase response factor)	BC000627	1.44
similar to H2A histone family, member A (H. sapiens)	BC001193	1.43
hypothetical protein FLJ13660 similar to CDK5 activator-binding protein C53	NM_025197	1.43
prelensin associated rhomboid-like protein	NM_018622	1.43
glucocorticoid receptor DNA binding factor 1	AU159484	1.43
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O	NM_020988	1.43
a disintegrin and metalloproteinase domain 10	NM_001110	1.43
thymidine kinase 1, soluble	NM_003258	1.43
programmed cell death 8 interacting protein	NM_013374	1.43
ubiquinol-cytochrome c reductase (8.4kD) subunit	NM_006830	1.43
zinc finger protein 313	NM_018683	1.43
chromobox homolog 4 (Pc class homolog, Drosophila)	NM_003655	1.43
Consensus includes gp:AV728528 /FEA=EST /DB_XREF=gi:10837947 /DB_XREF=est:AV728528 /CLONE=HTCBBDD04 /UG=Hs.62264 KIAA0937 protein	AV728526	1.43
KIAA0179 protein	AI400490	1.43
RAD1 homolog (S. pombe)	D79998	1.43
proliferating cell nuclear antigen	AF074717	1.43
NADH dehydrogenase (ubiquinone) Fe-S protein 2 (49kD) (NADH-coenzyme Q reductase)	NM_002592	1.43
mitogen-activated protein kinase 1	NM_004550	1.43
katanin p80 (WD40-containing) subunit B 1	NM_002745	1.43
MAGEF1 protein	NM_005886	1.43
Jun D proto-oncogene	NM_022149	1.43
nunt-related transcription factor 3	A1339541	1.43
CHK1 checkpoint homolog (S. pombe)	NM_004350	1.43
synapsin 10	NM_001274	1.43
soluble carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc) transporter), member 3	NM_003765	1.43
hypothetical protein FLJ11808	NM_012243	1.43
phosphoglycerate kinase 1	NM_024824	1.43
polyamine-modulated factor 1	NM_000291	1.43
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 7	NM_007221	1.43
N-myristoyltransferase 1	NM_012152	1.43
Consensus includes gp:AW573378 /FEA=EST /DB_XREF=gi:7246918 /DB_XREF=est:U1-HF-8L0-adc-e-05-0-U1.s1 /CLONE=IMAGE:3060969 /UG=Hs.6551 ATPase, H ⁺ trans	AF020500	1.43
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine polypeptide-N-acetylglucosaminyl transferase)	BF001665	1.43
glucosaminyl (N-acetyl) transferase 2, branching enzyme	L19659	1.43
v-abl Abelson murine leukemia viral oncogene homolog 1	NM_005157	1.43
IKK-related kinase epsilon; Inducible IkappaB kinase	NM_014002	1.43
methionine-tRNA synthetase	AA621558	1.43
RE1-silencing transcription factor	NM_005612	1.43
ubiquitin-conjugating enzyme HBUCE1	NM_015983	1.43
subtransferase family, cytosolic, 1C, member 2	NM_006588	1.43
adenylyate kinase 2	AL513611	1.43
Consensus includes gp:AI658004 /FEA=EST /DB_XREF=gi:5511820 /DB_XREF=est:wj69009.x1 /CLONE=IMAGE:2408057 /UG=Hs.284277 Homo sapiens immunoglobulin m	AI658004	1.43
hypothetical protein FLJ21986	NM_024913	1.43
reticulocalbin 2, EF-hand calcium binding domain	BC004892	1.43
yeast Sec31p homolog	AB020712	1.43
cholinergic receptor, nicotinic, alpha polypeptide 4	X89745	1.43
LGN protein	NM_013296	1.43
hypothetical protein MG20727	N32831	1.43
death effector domain-containing	AF064605	1.43
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	NM_002819	1.43
fragile X mental retardation, autosomal homolog 1	NM_005087	1.43
soluble carrier family 25 (carnitine/acylcarnitine translocase), member 20	BC001689	1.43
meningioma expressed antigen 6 (coiled-coil proline-rich)	U73682	1.43
hypothetical protein FLJ21062	NM_024788	1.43
glutaminase	AB020645	1.43
RNA binding motif protein 9	AW149379	1.43
stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4	AL136807	1.43
SET domain and mariner (transposase) fusion (fmm)	NM_006515	1.42

major histocompatibility complex, class I, J (pseudogene)	M60469	1.40
cytochrome P450, subfamily 1J (arachidonic acid epoxidase) polypeptide 2	NM_000775	1.40
blinervin reductase B (flavin reductase (NADPH))	NM_000713	1.40
chromosome 21 open reading frame 66	NM_016631	1.40
hypothetical protein FLJ14888	AF181382	1.40
entylin repeat and SOCS box-containing 1	AF055024	1.40
hypothetical protein FLJ10581	NM_016146	1.40
colony stimulating factor 1 (macrophage)	NM_000757	1.40
hydroxyacyl-Coenzyme A dehydrogenase/3-ketacyl-Coenzyme A thiolase/acyl-Coenzyme A hydratase (trifunctional protein), beta subunit	NM_000183	1.40
IQ motif containing GTPase activating protein 1	D29840	1.40
zinc finger protein 95 homolog (mouse)	BF196931	1.40
eml1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate)	BG475299	1.40
CASP8 and FADD-like apoptosis regulator	AF015451	1.40
YME1-like 1 (S. cerevisiae)	AF070656	1.40
elcC homolog 2 (E. coli)	NM_018127	1.40
RAM binding protein 18	NM_015024	1.40
LPAP for lysophosphatidic acid phosphatase	NM_016361	1.40
hypothetical protein DKFZP762M115	BC022477	1.40
calumenin	BF936365	1.40
ret finger protein	NM_006510	1.40
dishevelly/peptidase III	NM_005700	1.40
plakophilin 4	NM_003628	1.40
acyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase)	AW072302	1.40
ribosomal protein L37a	NM_000998	1.40
intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	A1608725	1.40
splicing factor 1	NM_004630	1.40
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 56/58kD, isoform 1 (Renal tubular acidosis with deafness)	NM_001692	1.40
cell division cycle 23C	NM_001790	1.40
novel putative protein similar to YIL091C yeast hypothetical 84 kD protein from SGA1-KTR7	N30910	1.40
CD64 antigen (leukocyte antigen)	NM_003874	1.40
polymerase (DNA-directed), alpha (70kD)	NM_002689	1.39
RNA polymerase I subunit	NM_004875	1.39
SET binding factor 1	U93181	1.39
receptor (TNFRSF)-interacting serine-threonine kinase 1	U50062	1.39
RAD23 homolog A (S. cerevisiae)	NM_005053	1.39
splicing factor 3b, subunit 1, 155kD	NM_012433	1.39
hypothetical protein FLJ11184	NM_018346	1.39
phosphatase and tensin homolog (mutated in multiple advanced cancers 1), pseudogene 1	AF023139	1.39
hypothetical protein FLJ10592	NM_022914	1.39
hypothetical protein 24638	BF439241	1.39
ab1-interactor 12 (SH3-containing protein)	NM_018147	1.39
proteasome (prosome, macropain) 26S subunit, non-ATPase, 12	BC002840	1.39
LJM protein (similar to rat protein kinase C-binding enigma)	AF260261	1.39
vacuolar protein sorting 18 (yeast)	A1446530	1.39
Consensus includes gb-M74089.1 (DEF-Human TB1 gene mRNA, 3 end. /FEA-mRNA/GEN=TB1/DB_XREF=gi:182400 /UG=Hs.75639 Human TB1 gene mRNA, 3 end	NM_008457	1.39
CCR4-NOT transcription complex, subunit 2	NM_022575	1.39
primase, polypeptide 1 (49kD)	M74089	1.39
PCTAIRE protein kinase 2	EG105204	1.39
vesicular endothelial growth factor	NM_000946	1.39
ceroid-lipofuscinosis, neuronal 5	NM_002395	1.39
G protein-coupled receptor 38	M27281	1.39
a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	NM_006493	1.39
tetrapsan 3	NM_001507	1.39
TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 32 kD	A1787833	1.39
H1 histone family, member 0	BC000704	1.39
KIAA0020 gene product	NM_003187	1.39
SH3 domain binding glutamic acid-rich protein like	Z97630	1.39
hypothetical protein FLJ10120	NM_014878	1.39
hypothetical protein FLJ23022	AL515318	1.39
Cluster Incl. AW001777:wd4f09.x1 Homo sapiens cDNA, 3' end (clone=IMAGE-2498233 /clone_end=3' (gb=AW001777 /gi=5848693 /ug=Hs.233841 /len=485	NM_018001	1.39
KIAA0019 protein	NM_025051	1.39
	AW001777	1.39
	AB020626	1.39

protein kinase, AMP-activated, beta 1 non-catalytic subunit
KIAA0456 protein
zinc finger protein 117 (HPE9)
RAP2A, member of RAS oncogene family
peptide-N-glycanase similar to yeast PNG1
sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)
KIAA0372 protein
Consensus includes gb:BG537255 /FEA=EST /DB_XREF=gi:13529117 /DB_XREF=est:602565318F1 /CLONE=IMAGE:4689748 /UG=Hs.82646 DnaJ (Hsp40) homolog, subla 8G537255
DEADH (Asp-Glu-Ala-Asp)His box polypeptide 3
general transcription factor II.1
neuronal PAS domain protein 2
DNFZP580F1916 protein
30 kDa protein
KIAA0076 gene product
uroporphyrinogen III synthase (congenital erythropoietic porphyria)
heparan sulfate 6-O-sulfotransferase
RNA binding motif, single stranded interacting protein 1
signal recognition particle 9kD
Consensus includes gb:AK023154.1 /DEF=Homo sapiens cDNA FLJ13092 (s. clone NT2RP002147, /FEA=mrna /DB_XREF=gi:10434948 /UG=Hs.172035 hypothetical prot AK023154
nuclear cap binding protein subunit 2, 20kD
LPS-induced TNF-alpha factor
filamin B, beta (actin binding protein 278)
myosin, light polypeptide 4, alkali; atrial; embryonic
APG3 autophagy 5-like (S. cerevisiae)
adducin 1 (alpha)
mitochondrial ribosomal protein L15
hypothetical protein FLJ21080
splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)
reqlum, apoptosis response zinc finger gene
desmin
F-box only protein 9
tyrosyl-tRNA synthetase
KIAA0365 gene product
chromosome 1 open reading frame 28
JAK binding protein
B-cell CLL/lymphoma 7A
hypothetical protein MGC3180
sorting nexin 6
hypothetical protein MGC10882
lamin B2
inositol polyphosphate-4-phosphatase, type I, 107kD
nuclear pore complex interacting protein
chromosome 5 open reading frame 4
cleavage and polyadenylation specific factor 4, 30kD subunit
autotransferase family, cytosolic, 1A, phenol-preferring, member 3
translocation protein 1
ribokinase
Consensus includes gb:AL008627 /DEF=Human DNA sequence from PAC 130G2 on chromosome 6p22.2-22.3. Contains ribosomal protein L29 pseudogene, ESTs and STS
ribonuclease P (14kD)
metastasis associated 1
protein phosphatase 1, regulatory (inhibitor) subunit 2
DEADH (Asp-Glu-Ala-Asp)His box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin)
hypothetical protein FLJ2427
tyrosylprotein sulfotransferase 2
foditin 1
anaphase-promoting complex subunit 10
proline oxidase homolog
transmembrane 7 superfamily member 2
carboxypeptidase D
DNFZP414N043 protein
chondrocytogen B1
eukaryotic translation initiation factor 3, subunit 7 (zeta, 66/67kD)
hypothetical protein FLJ20186

BC001007 1.38
AA742261 1.38
NM_024498 1.38
AI302106 1.38
NM_018297 1.38
AI743762 1.38
AI629033 1.38
8G537255 1.38
AF061327 1.38
AC004883 1.38
NM_002518 1.38
BE645501 1.38
NM_018447 1.38
NM_014780 1.38
NM_000375 1.38
NM_004807 1.38
NM_016637 1.38
NM_003133 1.38
AK023154 1.38
BC001255 1.38
NM_004862 1.38
AV12733 1.38
X58851 1.38
AK001889 1.38
BE898839 1.38
NM_014175 1.38
NM_022743 1.38
AI871457 1.38
NM_006268 1.38
NM_001927 1.38
AF176704 1.38
W27419 1.38
AV738039 1.38
AW192876 1.38
AI056051 1.38
NM_020993 1.38
NM_024041 1.38
NM_021249 1.38
BC004852 1.38
M94363 1.38
NM_004027 1.38
AA308653 1.38
H93077 1.38
NM_006693 1.38
U08032 1.38
U83239 1.38
NM_022128 1.38
AL008627 1.38
AV242755 1.38
BE561596 1.38
BF966540 1.38
NM_001357 1.38
BF129339 1.38
NM_003595 1.38
AF085357 1.38
NM_014885 1.38
AA074145 1.38
AF086304 1.38
NM_001304 1.38
8G538800 1.38
AW584011 1.38
NM_003753 1.38
NM_017703 1.38

ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) 1.38
 chromosome 9 open reading frame 7 1.38
 nuclear pore complex interacting protein 1.38
 KIAA1083 protein 1.38
 hypothetical protein MGC2454 1.38
 SRp25 nuclear protein 1.38
 proline rich 1 1.38
 nuclear cytochrome-rich protein 1.38
 nucleogen-activated protein kinase 1 1.38
 MCH33 mitochondrial maltenance deficient 3 (S. cerevisiae) 1.38
 hypothetical protein FLJ21977 1.38
 Bruton agammaglobulinemia tyrosine kinase 1.38
 RNA polymerase I transcription factor RRN3 1.38
 origin recognition complex, subunit 2-like (yeast) 1.38
 Consensus includes gb:AV700332 /FEA=EST /DB_XREF=gi:10302303 /DB_XREF=gi:10302303 /CLONE=GKCDIF.11 /UG=Hs.243501 Homo sapiens cDNA FLJ20738 fls, ck AV700332 1.38
 butyrate-induced transcript 1 1.38
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19KD, ASH1) 1.38
 CD3D antigen, delta polypeptide (T173 complex) 1.38
 protein tyrosine phosphatase, non-receptor type 6 1.38
 proenkephalin 1.38
 retinol x receptor interacting protein 1.38
 solute carrier family 35 (UDP-galactose transporter), member 2 1.38
 hypothetical protein FLJ20294 1.38
 KIAA0592 protein 1.38
 Consensus includes gb:AL080233.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586L111 (from clone DKFZp586L111) /FEA=mRNA /DB_XREF=gi:5262726 /UG=Hs.326580 H AL080233 1.38
 transcriptional adaptor 2 (ADA2 homolog, yeast)-like 1.38
 apoptosis-associated speck-like protein containing a CARD 1.38
 Bicaudal D homolog 1 (Drosophila) 1.38
 SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome) 1.37
 pre-B lymphocyte gene 3 1.37
 hypothetical protein FLJ10509 1.37
 nuclear VCP-like 1.37
 DC13 protein 1.37
 zinc finger RNA binding protein 1.37
 membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) 1.37
 activated RNA polymerase II transcription colicor 4 1.37
 HLA class II region expressed gene KE2 1.37
 karyopherin (importin) beta 1 1.37
 histone acetyltransferase 1.37
 poly(A) binding protein, cytoplasmic, pseudogene 3 1.37
 Consensus includes gb:T33069 /FEA=EST /DB_XREF=gi:615166 /DB_XREF=est:EST56576 /UG=Hs.7101 anaphase-promoting complex subunit 5 1.37
 hypothetical protein FLJ14007 1.37
 hypothetical protein FLJ10878 1.37
 nuclear pore complex interacting protein 1.37
 cornichon-like 1.37
 gb:NM_001514.2 /DEF=Homo sapiens general transcription factor IIB (GTF2B), mRNA. /FEA=mRNA (GEN=GTF2B) /PROD=general transcription factor IIB /DB_XREF=gi:1343 NM_001514 1.37
 Consensus includes gb:BG255188 /FEA=EST /DB_XREF=gi:12765004 /DB_XREF=est:8023697271 /CLONE=IMAGE:4477807 /UG=Hs.3343 phosphoglycerate dehydrogenase BG255188 1.37
 mitochondrial ribosomal protein S30 1.37
 hypothetical protein FLJ20258 1.37
 replication protein A1 (70KD) 1.37
 glucocorticoid modulatory element binding protein 1 1.37
 transmembrane 4 superfamily member 7 1.37
 trophoblast associated protein (tastin) 1.37
 cytosolic acyl coenzyme A thioester hydrolase 1.37
 KIAA0652 gene product 1.37
 NG23 protein 1.37
 neuroblastoma binding protein 8 1.37
 RAB28, member RAS oncogene family 1.37
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 1.37
 G52 gene 1.37
 uncharacterized bone marrow protein BM045 1.37
 MLN51 protein 1.37
 acyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) 1.37

52

forthead box O1A (nabdomyosarcoma)
 serine palmitoyltransferase, long chain base subunit 2
 RAB35, member RAS oncogene family
 NS1-associated protein 1
 inhibitor of DNA binding 2, dominant negative helix-loop-helix protein
 DNA segment, single copy, probe pH4 (transforming sequence, thyroid-1)
 general transcription factor IIIA
 carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase
 phosphoprotein enriched in astrocytes 15
 exosome component Rnp41
 inositol 1,4,5-trisphosphate receptor, type 3
 dual specificity phosphatase 1
 hypothetical protein FLJ11336
 methyl-CpG binding domain protein 3
 hypothetical protein FLJ22494
 hydromethylbilane synthase
 hypothetical protein R33729.1
 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)
 calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
 steroidogenic acute regulatory protein related
 dyenin, cytoplasmic, light polypeptide
 calcium phosphatase 2, regulatory subunit B (B56), gamma isoform
 protein phosphatase 2, regulatory subunit B (B56), gamma isoform
 TGF-beta-induced anti-apoptotic factor 1
 KIAA0365 gene product
 hypothetical protein FLJ21459
 acetyl-Coenzyme A transporter
 hypothetical protein FLJ10652
 bone morphogenetic protein 15
 leucine zipper transcription factor-like 1
 KIAA0683 gene product
 KIAA0876 protein
 Consensus includes gb:AI007083 /FEA=EST /DB_XREF=gi:6497611 /DB_XREF=est:PM-BT134-050499-650 /UG=Hs.124620 ESTs
 Consensus includes gb:AL133015.1 /IDEF=Homo sapiens mRNA, cDNA DKFZp434O2417 (from clone DKFZp434O2417); partial cds. /FEA=mRNA /GEN=DKFZp434O2417 IF AL133015
 amplified in osteosarcoma
 KIAA0794 protein
 similar to vaccinia virus HindIII K4L ORF
 methyl-CpG binding domain protein 1
 hypothetical protein FLJ20560
 splicing factor, arginine/serine-rich 6
 Consensus includes gb:AJ295618 /DEF=Homo sapiens FISH gene for putative ATPases, exons 1 and 2 and join CDS /FEA=mRNA /DB_XREF=gi:8506352 /UG=Hs.206521 Y AJ295618
 eriodine homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)
 zinc finger protein, subfamily 1A, 1 (Ikroos)
 gastrin-releasing peptide receptor
 solute carrier family 5 (sodium/glucose cotransporter), member 2
 hypothetical protein FLJ10335
 protein kinase, cAMP-dependent, regulatory, type II, beta
 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
 mitochondrial ribosomal protein S18A
 lysosomal-associated membrane protein 1
 SRY (sex determining region Y)-box 10
 solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 4
 transformer-2 alpha (htra-2 alpha)
 hypothetical protein FLJ22573
 KIAA0244 protein
 cholinergic receptor, nicotinic, alpha polypeptide 4
 TAK1-binding protein 2
 GABA(A) receptor-associated protein-like 2
 SSA protein SS-56
 leucine aminopeptidase
 high-mobility group (nonhistone chromosomal) protein 14
 ras homolog gene family, member C
 nuclear and coiled-body phosphoprotein 1
 UDP-Gal4betaGlcNAc 6-epimerase 1, 4- galactosyltransferase, polypeptide 5

potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3
 SNRNP assembly defective 1 homolog
 DKFZP566C134 protein
 KIAA0141 gene product
 hypothetical protein, expressed in osteoblast
 putative mitochondrial space protein 32.1
 programmed cell death 4 (neoplastic transformation inhibitor)
 insulin-degrading enzyme
 death-associated protein 6
 KIAA0930 protein
 methyltetrahydrofolate dehydrogenase (NADP+ dependent), methyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase
 serologically defined breast cancer antigen 64
 CGI-43 protein
 KIAA0153 protein
 hypothetical protein FLJ120411
 dolichyl-P-Glc:Man8GlcNAc2-PP-dolichylglucosyltransferase
 asparnylglicosaminidase
 purinergic receptor P2Y, G-protein coupled, 11
 hemochromatosis
 translocated promoter region (to activated MET oncogene)
 SHVSNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1
 testis-specific transcript, Y-linked 1
 phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase
 growth factor independent 1B (potential regulator of CDKN1A, translocated in CML)
 blast cell autoantigen 1 (69kD)
 gamma-aminobutyric acid (GABA) A receptor, alpha 5
 vesicle trafficking, beach and anchor containing
 transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase)
 CGI-02 protein
 Consensus includes gb:BE8986549 /FEA+EST /DB_XREF=gi:10369681 /DB_XREF=est:7164b02.x1 /CLONE=IMAGE:3299403 UG=Hs 44131 KIAA0974 protein
 apoptosis inhibitor
 melanoma antigen, family B, 4
 RAB3 GTPase-ACTIVATING PROTEIN
 tropomyosin 1 (alpha)
 upstream transcription factor 2, c-fos interacting
 GPAA1P anchor attachment protein 1 homolog (yeast)
 hypothetical protein similar to mouse aldolase reductase 6 (renal)
 tetraatropptide repeat domain 3
 fem-1 homolog b (C. elegans)
 DKFZP566L0724 protein
 KIAA1097 protein
 hypothetical protein FLJ10658
 gamma-glutamyltransferase-like 3
 Consensus includes gb:AK026977.1 /DEF=Homo sapiens cDNA: FLJ23324 fls, clone HEP12482, highly similar to HUMAHYOHCB Human nonmuscle myosin heavy chain-B (MN
 KIAA0418 gene product
 biased-like kinase 2
 U6 snRNA-associated Sm-like protein LSM8
 ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)
 hypothetical protein
 KIAA0901 protein
 deleted in lung and esophageal cancer 1
 hypothetical protein FLJ10422
 heterogeneous nuclear ribonucleoprotein A/B
 up-regulated by BCG-CWS
 hexosaminidase A (alpha polypeptide)
 hypothetical protein
 dihydrofolate reductase
 ZAP3 protein
 leukocyte receptor cluster (LRC) member 4
 potassium voltage-gated channel, shaker-related subfamily, beta member 1
 95 kDa rethroblastoma protein binding protein
 E2F transcription factor 1
 eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD)

NM_002249 1.36
 NM_006590 1.36
 AF161402 1.36
 NM_014773 1.36
 NM_006820 1.36
 AF050198 1.36
 NM_014456 1.36
 NM_004969 1.36
 NM_001350 1.36
 AK025608 1.36
 NM_005956 1.36
 AF091085 1.36
 NM_016011 1.36
 BF965437 1.36
 AB050442 1.36
 NM_013339 1.36
 NM_000027 1.36
 NM_002566 1.36
 AF115264 1.36
 NM_003292 1.36
 AL040633 1.36
 AF000990 1.36
 AF008655 1.36
 NM_004188 1.36
 NM_004968 1.36
 AF061785 1.36
 W60686 1.36
 BC003551 1.36
 NM_012123 1.36
 BE565549 1.36
 NM_021831 1.36
 NM_002367 1.36
 AK022494 1.36
 NM_000366 1.36
 NM_003367 1.36
 NM_003801 1.36
 AL096767 1.36
 AU131711 1.36
 AW138179 1.36
 AU158148 1.36
 AB029020 1.36
 NM_018248 1.36
 AL049709 1.36
 NM_002697 1.36
 NM_014631 1.36
 BF112255 1.35
 NM_016200 1.35
 AA602532 1.35
 NM_019083 1.35
 AB011173 1.35
 AB60341 1.35
 NM_018091 1.35
 NM_004499 1.35
 AL049963 1.35
 AL523158 1.35
 AF151022 1.35
 BC000192 1.35
 AB92986 1.35
 BC006309 1.35
 U16953 1.35
 AI379492 1.35
 NM_005225 1.35
 BE614908 1.35

hypothetical protein FLJ10774
Consensus includes gb:W84525 /FEA=EST /DB_XREF=gi:1395637 /DB_XREF=est:zd9g09.s1 /CLONE=IMAGE:356800 /UG=Hs.7949 DKFZP56862420 protein
G protein-binding protein CRFG
heat shock 70kD protein 4
Smey homolog, Y chromosome (mouse)
tumor protein, translationally-controlled 1
protein kinase C, δ
DEAD(H) (Asp-Glu-Ala-Asp-His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukaphysin)
phosphoglucosylase 1
CGI-203 protein
Consensus includes gb:A1348006 /FEA=EST /DB_XREF=gi:4085212 /DB_XREF=est:qp61b05.x1 /CLONE=IMAGE:1927473 /UG=Hs.469 succinate dehydrogenase complex, s
NICE-5 protein
senh1/SUMO-specific protease 3
cyclin-dependent kinase 9 (CDC2-related kinase)
protein phosphatase 1, regulatory (inhibitor) subunit 2
hypothetical protein DT1P1A10
Bardet-Biedl syndrome 4
protein tyrosine phosphatase type IVA, member 2
hypothetical protein
Sp-B transcription factor (Spi-1/PU.1 related)
zinc finger protein 45 (z Kruppel-associated box (KRAB) domain polypeptide)
heat shock factor binding protein 1
hypothetical protein FLJ22822
Cluster Inc. AI982878:wd24c06.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2508394 /clone_end=3 /gb=AI962879 /gi=5755592 /ug=Hs.81920 /len=430
H2B histone family, member L
Consensus includes gb:BGQ31677 /FEA=EST /DB_XREF=gi:12422203 /DB_XREF=est:602300830F1 /CLONE=IMAGE:4402326 /UG=Hs.58210 ESTs
hypothetical protein FLJ22237
hypothetical protein FLJ20073
solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5
potassium channel modulatory factor
ubiquitin specific protease 13 (isopeptidase T-3)
SAR1 protein
serine/threonine kinase 24 (STE20 homolog, yeast)
MGIM4 mitochondriome maintenance deficient 4 (S. cerevisiae)
hypothetical protein FLJ11937
hypothetical protein FLJ11937
Onu1 (Hsp40) homolog, subfamily A, member 3
Consensus includes gb:AI798823 /FEA=EST /DB_XREF=gi:3384295 /DB_XREF=est:w693b05.x1 /CLONE=IMAGE:2348625 /UG=Hs.1050 pleckstrin homology, Sec7 and coiled coil
vev 2 oncogene
hypothetical protein MGC4022
amyloid beta (A4) precursor protein-binding, family B, member 1 (Fe65)
lactate dehydrogenase B
methyl CpG binding protein 2 (Rett syndrome)
ectonucleoside triphosphate diphosphohydrolase 6 (putative function)
ribosomal protein S4, X-linked
methyl-CpG binding domain protein 4
ubiquitin specific protease 14 (IRNA-quinine transglycosylase)
protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta)
protein tyrosine kinase 9
nuclear protein ANK1
peroxisomal acyl-CoA thioesterase 1
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2
1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)
potassium voltage-gated channel, shaker-related subfamily, member 6
hypothetical protein
hypothetical protein
ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) 42kD
tyrosyl-DNA phosphodiesterase
regulator of Fas-induced apoptosis
ATPase, Na⁺/K⁺ transporting, alpha 1 polypeptide
hypothetical protein from EUROIMAGE 1987170
glutathione transferase zeta 1 (maleylacetoacetate isomerase)

SET domain, bifurcated 1
 Consensus includes gb:AL024509 /DEF=Human DNA sequence from clone 52P213 on chromosome 6p21.31-22.3. Contains a 60S Ribosomal Protein L21 pseudogene and a hypoxia-inducible factor 1, alpha subunit inhibitor
 metallothionein 1G
 aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase)
 immunoglobulin kappa variable 1J015-118
 Consensus includes gb:A071147 /FEA=EST /DB_XREF=gi:4651478 /DB_XREF=IMAGE:2297218 /UG=Hs.81360 Homo sapiens cDNA: FLJ121927 fr. A071147
 plectatrin
 ribosomal protein S7
 pituitary tumor-transforming 1
 Consensus includes gb:AA650558 /FEA=EST /DB_XREF=gi:2577886 /DB_XREF=IMAGE:1192002 /UG=Hs.325202 ESTs, highly similar to GBAS_H1, AA650558
 solute carrier family 2 (facilitated glucose transporter), member 1
 hypothetical protein FLJ21839
 cleavage stimulation factor, 3' pre-RNA, subunit 3, 77ND
 tetranucleotide repeat domain 4
 RAB2, member RAS oncogene family
 erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked)
 inhibitor of DNA binding 4, dominant negative helix-loop-helix protein
 hypothetical protein FLJ20827
 fibroblast growth factor 5
 Consensus includes gb:AX022663.1 /DEF=Homo sapiens cDNA FLJ12601 fts, clone NT2RM4001414, /FEA=mRNA /DB_XREF=gi:10434187 /UG=Hs.288549 Homo sapiens c AK022663
 lamin B1
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5 (16kD, SGDh)
 chromodomain protein, Y chromosome-like
 iniquitos homeobox protein 5
 nuclear factor (erythroid-derived 2)-like 1
 aryl hydrocarbon receptor nuclear translocator
 Consensus includes gb:AL133173 /DEF=Human DNA sequence from clone RP11-453N3 on chromosome 10 Contains the 3 end of a gene similar to ABCD1 (ATP-binding cassette AL133173)
 hypothetical protein MGC11308
 carbonic anhydrase II
 mannosidase, alpha, class 2B, member 1
 ribosomal protein L34
 Consensus includes gb:AL109622 /DEF=Human DNA sequence from clone RP3-526F5 on chromosome Xq26.3-28. Contains the gene for a novel protein similar to UBE2N (ut S55735)
 immunoglobulin heavy constant mu
 ras-like protein
 growth factor receptor-bound protein 2
 KIAA1449 protein
 ubiquitin-conjugating enzyme E2 variant 1
 FLN29 gene product
 Rho guanine exchange factor (GEF) 12
 Cluster Ind. N32782:yy91903.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:239638 /done_end=3 /gb=N32782 /gi=1153181 /ug=Hs.13138 /len=668
 solute carrier family 35 (CMP-sialic acid transporter), member 1
 peroxisomal biogenesis factor 3
 sine oculis homeobox homolog 2 (Drosophila)
 hypothetical protein FLJ12525
 second mitochondria-derived activator of caspase
 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit b, isoform 1
 putative zinc finger protein from EUROMIMAGE 566589
 aspartyl aminopeptidase
 hypothetical protein FLJ12442
 collagen
 Consensus includes gb:AL049987.1 /DEF=Homo sapiens mRNA: cDNA DKFZ564F112 (from clone DKFZ564F112), /FEA=mRNA /DB_XREF=gi:4684238 /UG=Hs.166361 H AL049987
 hypothetical protein FLJ22477
 glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)
 hypothetical protein FLJ12847
 HSPC166 protein
 similar to Caenorhabditis elegans protein C42C1.9
 gogli associated, gamma adiplin ear containing, ARF binding protein 2
 protocadherin beta 17 pseudogene
 hypothetical protein FLJ20550
 general transcription factor IIF, polypeptide 2 (44kD subunit)
 breast cancer anti-estrogen resistance 3

Consensus includes gb:AL359954 /DEF=Human DNA sequence from clone RP11-318P23 on chromosome 20 Contains a TAR DNA-binding protein pseudogene, ESTs, STSs. AL359954
Consensus includes gb:AF070537.1 /DEF=Homo sapiens clone 24606 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387894 /UG=Hs.17481 Homo sapiens clone 24606 mRt AF070537
squalene epoxidase
CDP-diclycydol synthase (phosphatidate cytidylyltransferase) 2
hepatocyte growth factor-regulated tyrosine kinase substrate
AFG3 ATPase family gene 3-like 2 (yeast)
retinoblastoma binding protein 4
syntaxin 7
diaminoguan
hypothetical protein FLJ10462
growth differentiation factor 1
Cluster Ind. AA601208:nt14c12.1 Homo sapiens cDNA. 3' end /clone=IMAGE:1106682 /clone_end=3 /gb=AA601208 /gi=2434833 /ug=Hs.220255 /len=488
early developmental regulator 2 (homolog of polyomaelic 2)
hypothetical protein FLJ12528
gb:NM_005703.2 /DEF=Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA. /FEA=CDS /GEN=UREB1 /PROD=upstream regulatory element binding protein 1 (UREB1), mRNA. /FEA=mRNA /DB_XREF=gi:4884400 /UG=Hs.22598 /AL050185
tridecaolide repeat containing 3
BUP protein
hypothetical protein FLJ20360
hypothetical protein FLJ22584
creatine kinase, muscle
cell division cycle 42 (GTP binding protein, 25kD)
active BCR-related gene
hypothetical protein DKFZp782A227
transposon-derived Buster3 transposase-like
HSPC022 protein
hypothetical protein FLJ11506
ubiquitin specific protease 16
membrane-associated tyrosine- and threonine-specific cdc2-inhibitory kinase
hypothetical protein FLJ11848
interferon, alpha 1
thymosin, beta 4, Y chromosome
nuclear phosphoprotein similar to S. cerevisiae PWP1
heparan sulfate 2-O-sulfotransferase
insulin-degrading enzyme
ribosomal protein L38
ribosomal protein L13
hypothetical protein FLJ13868
DKFZP564O062 protein
hypothetical protein
interleukin 12 receptor, beta 1
vimentin
ubiquitin-conjugating enzyme E2 variant 1
FliJ homolog 1 (E. coli)
zhedrin
dynactin 2 (p50)
phospholipase C, beta 2
ION3 protein
NY-REN-24 antigen
cytochrome b-561
ras homolog gene family, member
phosphatidylinositol glycan, class N
E2F transcription factor 2
chromosome 5 open reading frame 8
Consensus includes gb:AL050185.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586A0423 (from clone DKFZp586A0423). /FEA=mRNA /DB_XREF=gi:4884400 /UG=Hs.22598 /AL050185
putative DNA binding protein
ribosomal protein S2
centaurin, beta 1
myosin, light polypeptide, regulatory, non-sarcomeric (20kD)
KIAA0385 gene product
amyloid beta (A4) precursor protein-binding, family B, member 2 (Fe65-like)
TNFRSF1A-associated via death domain
interferon gamma receptor 1
centaurin, alpha 1

poliovirus receptor-related 1 (herpesvirus entry mediator C; nectin)
 H3 histone, family 3B (H3.3B)
 KIAA0998 protein
 chromosome 11 open reading frame 23
 zinc finger protein, 103 homolog (mouse)
 hypothetical protein FLJ21865
 RNA helicase family
 nuclear antigen Sp100
 fragile X mental retardation 2
 AHNK nucleoprotein (desmoyotin)
 hypothetical protein FLJ20045
 Consensus includes gb:AU154486 /FEA=EST /DB_XREF=gi:11016007 /DB_XREF=est:AU154486 /CLONE=NT2RP4001238 /UG=Hs.119023 SMC2 (structural maintenance of Nt protein 2)
 H3 histone family, member 1
 hypothetical protein dJ45N24.2.1
 hypothetical protein FLJ22246
 RNA guanylyltransferase and 5'-phosphatase
 integral membrane protein 3
 copine 1
 thyroid hormone receptor interactor 4
 vesicular protein sorting 45B (yeast)
 IGF-II mRNA-binding protein 3
 ubiquitin-conjugating enzyme E2L 3
 hypothetical protein DKFZ6564D0478
 hypothetical protein FLJ10583
 encephelin
 ribosomal protein L31
 TERF1 (TRF1)-interacting nuclear factor 2
 Ctip1-interacting zinc finger protein
 proteasome (prosome, macropain) 26S subunit, ATPase, 2
 oysteroid binding protein-like 3
 Consensus includes gb:BF445142 /FEA=EST /DB_XREF=gi:11510203 /DB_XREF=est:na021b05.x1 /CLONE=IMAGE:3386081 /UG=Hs.797 nuclear transcription factor Y, alpha
 Consensus includes gb:AK025343.1 /DEF=Homo sapiens cDNA: FLJ21690 fta, clone COL08538, /FEA=mRNA /DB_XREF=gi:10437841 /UG=Hs.306802 Homo sapiens cDNA: AK022874
 tumor protein, translationally-controlled 1
 calpain 10
 CLIP-associated protein 2
 BRCA1 associated protein-1 (ubiquitin carboxy-terminal hydrolase)
 tubulin-specific chaperone 8
 chromosome 11 hypothetical protein ORF4
 splicing factor 3b, subunit 1, 155kD
 hypothetical protein FLJ20721
 chorionic somatomammotropin hormone 1 (placental lactogen)
 paraneoplastic antigen MA1
 protein-rich protein with nuclear targeting signal
 hypothetical protein DKFZp434E1723
 STIP1 homology and U-Box containing protein 1
 calcitriol/calmodulin-dependent protein kinase (CaM kinase) II beta
 BCL2-associated X protein
 low density lipoprotein receptor-related protein 8, apolipoprotein E receptor
 KIAA0690 protein
 zinc finger protein 277
 hypothetical protein FLJ13910
 UBX domain-containing 2
 protease, serine, 22
 ubiquitin specific protease 4 (proto-oncogene)
 KIAA0978 protein
 putative protein kinase NY-REN-64 antigen
 hypothetical protein DKFZp434K0920
 adaptor-related protein complex 4, sigma 1 subunit
 muS5 homolog 6 (E. coli)
 KIAA0372 gene product
 tripartite motif-containing 36

NM_002855 1.32
 AW138159 1.32
 AB023215 1.32
 NM_018312 1.32
 NM_005667 1.32
 AL044087 1.32
 AA156961 1.32
 NM_003113 1.32
 NM_002025 1.32
 BG287862 1.32
 BC002894 1.32
 NM_020202 1.32
 NM_021016 1.32
 AF247168 1.32
 NM_025232 1.32
 AB009023 1.32
 NM_030926 1.32
 NM_003915 1.32
 NM_016213 1.32
 AF165513 1.32
 AL023775 1.32
 AI819709 1.32
 AL136883 1.32
 NM_018148 1.32
 NM_001988 1.32
 NM_000993 1.32
 NM_012481 1.32
 NM_012127 1.32
 NM_002803 1.32
 AI202969 1.32
 BF445142 1.32
 AK025343 1.32
 AK022874 1.32
 AL565449 1.32
 NM_023089 1.32
 AI741784 1.32
 NM_004656 1.32
 NM_004607 1.32
 AA813184 1.32
 AI739389 1.32
 NM_017941 1.32
 NM_022641 1.32
 NM_006029 1.32
 AF279899 1.32
 AI743331 1.32
 NM_005861 1.32
 AF112472 1.32
 U19599 1.32
 NM_004631 1.32
 AK021460 1.32
 NM_021994 1.32
 BF671894 1.32
 AU149808 1.32
 NM_006025 1.32
 AI646043 1.32
 BE256900 1.32
 NM_016123 1.32
 NM_017590 1.32
 AF155159 1.32
 NM_000179 1.32
 BE566023 1.32
 NM_018700 1.32

3-phosphoinositide dependent protein kinase-1	AF017995	1.31
DKFZP586I2233 protein	BC002857	1.32
prinsase, polypeptide 2A (58KD)	NM_000947	1.32
Consensus includes gb:A7007128.1 IDEF=Homo sapiens clone CDABP0028 mRNA sequence. /FEA=mRNA /DB_XREF=ggi:596024 /JUG=Hs.24435 Homo sapiens clone A7007128	AW07128	1.32
KIA00692 protein	AW274877	1.32
myobularin related protein 2	U56033	1.32
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha	A0187167	1.32
Consensus includes gb:BF057059 /FEA=EST /DB_XREF=ggi:10810955 /DB_XREF=est:7k15c07.x1 /CLONE=IMAGE:3444468 /JUG=Hs.12421 hypothetical protein	BF057059	1.32
microtubule-associated proteins 1A1B light chain 3	BE593893	1.32
sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C	NM_030913	1.32
8-adenosylhomocysteine hydrolase-like 1	AA479488	1.32
v-ea erythroblastosis virus E28 oncogene homolog 2 (avian)	NM_005239	1.32
kyurenine 3-monooxygenase (lynurinine 3-hydroxylase)	A0174145	1.32
bone marrow stromal cell antigen 1	NM_004334	1.32
Consensus includes gb:BG429214 /FEA=EST /DB_XREF=ggi:13335720 /DB_XREF=est:602498083F1 /CLONE=IMAGE:4611836 /JUG=Hs.258144 zinc finger protein 33b (K0X	BG429214	1.32
biphenyl hydrolase-like (serine hydrolase; breast epithelial mucin-associated antigen)	NM_004332	1.32
hypothetical protein	NM_017446	1.32
zinc finger protein 143 (clone pHZ-1)	AW182015	1.32
upstream transcription factor 2, c-fos interacting	X90824	1.32
hypothetical protein MGC33067	NM_024295	1.32
hypothetical protein MGC2478	NM_024298	1.32
hypothetical protein FLJ20452	BE313890	1.32
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D	NM_006378	1.32
Consensus includes gb:AK023177.1 IDEF=Homo sapiens cDNA: FLJ21524 flg, clone COL05921. /FEA=mRNA /DB_XREF=ggi:10437640 /JUG=Hs.308778 Homo sapiens cDNA: AK023177	NM_024894	1.32
hypothetical protein FLJ12595	NM_024894	1.32
Consensus includes gb:BE338424 /FEA=EST /DB_XREF=ggi:87670689 /DB_XREF=est:601068256F1 /CLONE=IMAGE:3454693 /JUG=Hs.288283 Homo sapiens cDNA: FLJ2231	BE538424	1.32
translocase of inner mitochondrial membrane 44 homolog (yeast)	AF076030	1.32
pumilio homolog 1 (Drosophila)	BE670915	1.31
nuclear receptor coactivator 4	AL162047	1.31
mult. homolog 1, colon cancer, nonpolyposis type 2 (E. coli)	NM_000249	1.31
Sec23-interacting protein p125	AK021846	1.31
hypothetical protein FLJ20608	NM_017900	1.31
G protein-coupled receptor 27	NM_018971	1.31
klornate 2-sulfatase (Hunter syndrome)	NM_000202	1.31
tubulin, beta polypeptide	NM_001089	1.31
tumor rejection antigen (gp96) 1	AI582238	1.31
SKI-interacting protein	NM_012245	1.31
G protein-coupled receptor 43	NM_005308	1.31
endocrine regulator	NM_014345	1.31
VRK3 for vacuola related kinase 3	BF062866	1.31
protein phosphatase 4, regulatory subunit 1	NM_005134	1.31
Consensus includes gb:AW128846 /FEA=EST /DB_XREF=ggi:8118784 /DB_XREF=est:xd4806.x1 /CLONE=IMAGE:2598835 /JUG=Hs.24437 ESTs. Weakly similar to neurona	AW128846	1.31
hypothetical protein FLJ12009	NM_018360	1.31
Consensus includes gb:BC004409.1 IDEF=Homo sapiens, clone IMAGE:3638994, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3638994) /DB_XREF	BC004409	1.31
small nuclear ribonucleoprotein D3 polypeptide (18KD)	NM_004175	1.31
KIA00948 protein	AW991219	1.31
RAB48, member RAS oncogene family	NM_016154	1.31
RAP1, GTP-GDP dissociation stimulator 1	BC001851	1.31
hypothetical protein MGC2875	NM_024073	1.31
KIA00810 protein	BE615699	1.31
protein tyrosine phosphatase type IVA, member 2	AF208850	1.31
salicase 3 (membrane salicase)	AK022450	1.31
ubiquitin C-terminal hydrolase UCH37	NM_015984	1.31
glutamate-cysteine ligase, catalytic subunit	BF676880	1.31
C2H2 (Kruppel-type) zinc finger protein	AF159587	1.31
stannin	AF070673	1.31
prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)	S38219	1.31
PAS-serine/threonine kinase	D50925	1.31
5-hydroxytryptamine (serotonin) receptor 2C	NM1778	1.31
regulator of G-protein signalling 14	AF037194	1.31
paired-like homeodomain transcription factor 1	U70370	1.31
PAU-1 mRNA-binding protein	AF131807	1.31
axosomal protein L24	AI560573	1.31

SRY (sex determining region Y)-box 4	A089477	1.31
perlecanin 2 (kndrin)	NM_006031	1.31
SBB126 protein	NM_018946	1.31
calpastatin	AF327443	1.31
cleavage and polyadenylation specific factor 6, 68kD subunit	AU149387	1.31
parilin	D86862	1.31
hypothetical protein FLJ12686	NM_024595	1.31
2N10 homolog, centromeres/kinetochore protein (Drosophila)	NM_004724	1.31
hypothetical protein FLJ10057	NM_017684	1.31
hypothetical protein FLJ21657	NM_022483	1.31
G protein-coupled receptor 30	U63917	1.31
vacuolar protein sorting 35 (yeast)	NM_018206	1.31
solute carrier family 18 (vesicular monoamine), member 2	AZ69290	1.31
acrosomal vesicle protein 1	NM_020110	1.31
ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit (oligomycin sensitivity conferring protein)	S77356	1.31
cytochrome c oxidase subunit Va	NM_004255	1.31
PPTAIRE protein kinase 1	NM_012395	1.31
EGF-like domain, multiple 3	AL134003	1.31
hypothetical protein FLJ10280	NM_018047	1.31
heat shock transcription factor 4	AB029348	1.31
adenylyl cyclase-associated protein	AA806142	1.31
KIAA0876 protein	NM_015043	1.31
KIAA0873 protein	AB014573	1.31
mudS homolog 3 (E. coli)	NM_002439	1.31
DKFZP564J102 protein	AL060065	1.31
KIAA1232 protein	AB033056	1.31
zinc finger protein 212	NM_012256	1.31
Consensus includes gb:A1721229 /FEA=EST /DB_XREF=gi:5038485 /DB_XREF=est:as68c10.x1 /CLONE=IMAGE:2333874 /UG=Hs.326456 hypothetical protein FLJ20030	A1721229	1.31
tumoral hyaluronidase	NM_000143	1.31
ferredoxin 1	NM_004109	1.31
transcriptional co-activator with POZ-binding motif (TAZ)	BF674349	1.31
squamous cell carcinoma antigen recognised by T cells 3	AW173076	1.31
hypothetical protein MDS018	NM_021823	1.31
exosome component Rps41	AIS71288	1.31
glucocorticoid receptor DNA binding factor 1	AB070100	1.31
gb:AB056895.1 /DEF=Homo sapiens GHRHRp sv mRNA for growth hormone releasing hormone receptor pituitary splice variant, complete cds. /FEA=mrna /GEN=HHRp	NM_005895	1.31
lactotransferrin	NM_002343	1.31
tripartite motif-containing 28	NM_005762	1.31
Cluster Ind. W07773:zb03g04.1 Homo sapiens cDNA, 5' end /clone=IMAGE:301014 /clone_end=5' gb=W07773 /gi=1281785 /ug=Hs.9739 /len=533	W07773	1.31
HEMK homolog 7kb	AF131220	1.31
cholea acanthocytosis	AB023203	1.31
hypothetical protein MGC3121	AA161130	1.31
olfactory receptor, family 2, subfamily 5, member 2	NM_019897	1.31
uncharacterized hypothalamus protein HARP11	AK022248	1.30
KIAA0887 protein	NM_014938	1.30
CGI-148 protein	NM_016076	1.30
gene with multiple splice variants near HD locus on 4p16.3	NM_003704	1.30
MAOS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	AL536517	1.30
Consensus includes gb:AL390057 /DEF=Human DNA sequence from clone RP11-51N22 on chromosome 13 Contains ESTs, STSs and GSSs. Contains an HNRPA1 (heterogr	AL390057	1.30
secreted luteal-related protein 1	A132407	1.30
hypothetical protein FLJ10315	NM_018056	1.30
phosphatidylinositol binding diaphanin assembly protein	AV721177	1.30
CD19 antigen	NM_001770	1.30
vesicle-associated membrane protein 4	NM_003762	1.30
MZ7630 Human 26S ribosomal RNA gene, complete cds (.5, .M, .3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	MZ7630	1.30
suppressor of var1.3-like 1 (S. cerevisiae)	NM_003171	1.30
serine/threonine kinase 11 (Peutz-Jeghers syndrome)	AF036625	1.30
synovial sarcoma, X breakpoint 2	AU152068	1.30
KIAA0332 protein	NM_018933	1.30
procadherin beta 13	AV756141	1.30
colony stimulating factor 2 receptor, beta, low-affinity (granulocyte-macrophage)	M34356	1.30
cAMP responsive element binding protein 1	NM_019053	1.30
chromosome 2 open reading frame 2		

Consensus includes gb:J04742 /DEF=Human autonomous replicating sequence H1 (ARSH1) /FEA=CDS /DB_XREF=gi:179028 /UG=Hs.247345 Human autonomous replicatin J04742
KJAA0090 protein
Consensus includes gb:AK002078.1 /DEF=Homo sapiens cDNA FLJ11214 fs, clone PLACE1007990. /FEA=mRNA /DB_XREF=gi:7023738 /UG=Hs.281616 Homo sapiens cD AK002078
DNFZ2564C196 protein
inositol polyphosphate phosphatase-like 1
glycosylphosphatidylinositol specific phospholipase D1
hypotheical protein FLJ22855
8TB (PO2) domain containing 1
hypotheical protein FLJ20343
Consensus includes gb:M222005 /DEF=Human interleukin 2 gene, clone pATlacIL-2C2TT, complete cds, clone pATlacIL-2C2TT /FEA=CDS /DB_XREF=gi:186300 /UG=Hs.2471422005
zinc finger protein 36, C3H type-like 1
nuclear transport factor 2 (placental protein 15)
hypotheical protein MGC2483
zinc finger protein 278
fodilin 1
HSPC035 protein
Consensus includes gb:AL109942 /DEF=Human DNA sequence from clone RP3-47J18 on chromosome 6p25.3-26 Contains 3 end of the MAP3K4 (mitogen-activated protein AL109942
Cluster loc. A424872 /g38110.11 Homo sapiens cDNA, 3 and clone IMAGE:2111083 /clone_end=3 /gb=A424872 /g=4270895 /ug=Hs.159375 /len=443
DEADH (Asp-Glu-Ala-Asp-His) box polypeptide 10 (RNA helicase)
p6gi associated, gamma adspilin ear containing, ARP binding protein 2
hypotheical protein MGC355
Cluster loc. AW028481 /w14604.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:990913 /clone_end=3 /gb=AW028481 /g=5880011 /ug=Hs.48396 /len=486
alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I; glycolicaciduria; serine-pyruvate aminotransferase)
origin recognition complex, subunit 1-like (yeast)
Rab9 effector p40
phosphofructokinase, liver
zinc finger protein 16 (KIX 8)
nucleix (nucleoside diphosphate linked moiety X)-type motif 2
metallothionein IV
antigen identified by monoclonal antibody Ki-67
arsenate resistance protein ARS2
RAS guanyl releasing protein 2 (calcium and DAG-regulated)
glucose-6-phosphate dehydrogenase
KIAA0370 protein
hypotheical protein FLJ20783
nuclear transcription factor Y, alpha
prostate epithelium-specific Ets transcription factor
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)
ribosomal protein L7
hypotheical protein FLJ21840
Rho GTPase activating protein 1
hypotheical protein similar to preferentially expressed antigen of melanoma
solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulatory factor 1
CGI-30 protein
glycerol-3-phosphate dehydrogenase 2 (mitochondrial)
lysine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide
Consensus includes gb:L24521.1 /DEF=Human (transformation-related protein mRNA, 3 end, /FEA=mRNA /PROQ=transformation-related protein /DB_XREF=gi:403459 /UG=H L24521
nombold, veitid-like 1 (Drosophila)
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4
KIAA0046 protein
nuclear receptor co-repressor 2
chloride channel, calcium activated, family member 4
mannosidase, alpha, class 1B, member 1
heparin-binding growth factor binding protein
KIAA0063 gene product
fumarate hydratase
sorting nexin 11
DHHC1 protein
serine/threonine kinase 18
v-myo myeloblastosis viral oncogene homolog (avian)-like 1
protein tyrosine kinase 2 beta
small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen)
acidic protein rich in leucines

folate receptor 3 (gamma)	NM_000804	1.29
bedin 1 (coiled-coil, myosin-like BCL2 interacting protein)	NM_003786	1.29
hypothetical protein MGC4825	AF061264	1.29
dyxerlin, limb girdle muscular dystrophy 2B (autosomal recessive)	NM_003494	1.29
patently expressed 10	AL582836	1.29
surfeit 1	NM_003172	1.29
regulator of G-protein signalling 16	BF304996	1.29
regucalcin (senescence marker protein-30)	D31815	1.29
fatty-acid-Coenzyme A ligase, long-chain 2	NM_021122	1.29
granulin	BC000324	1.29
similar to yeast Upt3, variant B	NM_023010	1.29
gamma-glutamyl carboxylase	BE326952	1.29
OnaJ (Hsp40) homolog, subfamily C, member 4	NM_005528	1.29
Interleukin-1 receptor-associated kinase 1	NM_001569	1.29
hypothetical protein FLJ10719	BG403815	1.29
aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (3-alpha)-hydroxysteroid dehydrogenase)	S68290	1.29
rTS beta protein	NM_017512	1.29
transducin (beta)-like 2	AI608789	1.29
pre-B-cell colony-enhancing factor	NM_005746	1.29
solute carrier family 28 (nucleoside transporters), member 1	AF078117	1.29
karyopherin (importin) beta 1	AA861608	1.29
selenoprotein X, 1	NM_016332	1.29
hypothetical protein FLJ10688	AF062733	1.29
natural killer-tumor recognition sequence	AI688640	1.29
EH-domain containing 1	AF001434	1.29
filamin B, beta (actin binding protein 278)	M62994	1.29
RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)	NM_002875	1.29
LamC (bacterial lamibiotic synthetase component C)-like 2	NM_018697	1.29
v-naf musculoskeletal fibrosarcoma oncogene homolog F (avian)	NM_012323	1.29
protein kinase C, alpha	NM_002737	1.29
tumor necrosis factor, alpha-induced protein 3	NM_006290	1.29
mannosyl (alpha-1,6)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	NM_006290	1.29
Consensus includes gb:A149235 /FEA=EST /DB_XREF=gi:3678004 /DB_XREF=est:ac70f11.x1 /CLONE=IMAGE:1714989 /UG=Hs.24064 Homo sapiens mRNA; cDNA DKFZ	AW103285	1.29
KUAA0802 protein	AB018345	1.29
phosphochukinase, liver	NM_002626	1.29
gb:BC001685.1 /DEF=Homo sapiens, Similar to cadherin 1, type 1, E-cadherin (epithelial), done MGC:1151, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cadherin 1	NM_002626	1.29
hypothetical protein d434014.3	BC001865	1.29
Consensus includes gb:BF037823 /FEA=EST /DB_XREF=gi:10746209 /DB_XREF=est:601481994F1 /CLONE=IMAGE:3865487 /UG=Hs.9883 OnaJ (Hsp40) homolog, subfam	NM_025228	1.29
hypothetical protein FLJ13096	NM_025000	1.29
serine/arginine repetitive matrix 2	AI655799	1.29
biliverdin reductase A	AA740186	1.29
dystroglycan 1 (dystrophin-associated glycoprotein 1)	NM_004393	1.29
N-acetylneuraminic acid phosphate synthase; sialic acid synthase	NM_018946	1.29
alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae)	U09820	1.29
survival of motor neuron 1, telomeric	NM_000344	1.29
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)	AU150748	1.29
tripartite motif-containing 26	NM_003449	1.29
transcription factor 8 (represses interleukin 2 expression)	U12170	1.29
proteasome (prosome, multicatalytic) subunit, alpha type, 3	NM_002788	1.29
solute carrier family 7 (cationic amino acid transporter, y ⁺ system), member 8	AL365347	1.29
ALEX1 protein	NM_016608	1.29
leukocyte-IRNA synthetase	NM_013417	1.29
KUAA0908 protein	AA502912	1.29
mitochondrial ribosomal protein S31	AI937794	1.29
NAD, mothers against decapentaplegic homolog 2 (Drosophila)	NM_005901	1.29
cyclin M1	NM_020348	1.29
transcription elongation factor A (SII), 1	X57198	1.29
KUAA0737 gene product,	BE783632	1.29
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	U80918	1.29
ER to nucleus signalling 1	NM_001433	1.29
CDK3 antigen (melanoma 1 antigen)	NM_001780	1.29
complement component 8, beta polypeptide	NM_000066	1.29
angiodotin-3	NM_015985	1.29

Consensus includes gb:XB4340.1 /DEF=H.sapiens mRNA (or Ig light chain, variable region (ID:CLL001VL)) /FEA=mRNA /GEN=lg VL gene /PROD=immunoglobulin light chain X84340 1.29
 Hox and leucine-rich repeat protein 5 AF157323 1.29
 rth endonuclease III-like 1 (E. coli) AC005600 1.29
 H4 histone family, member G NM_003542 1.29
 Ras homolog enriched in brain 2 BF593727 1.28
 myobulin related protein 4 AB014547 1.28
 golgi associated, gamma adaptin ear containing, ARF binding protein 3 AV008018 1.28
 breast carcinoma amplified sequence 2 NM_005872 1.28
 SET translocation (myeloid leukemia-associated) NM_003011 1.28
 HUS1 checkpoint homolog (S. pombe) NM_004507 1.28
 transient receptor potential channel 1 U31110 1.28
 ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function) AW194947 1.28
 hypothetical protein MGC12335 AL022724 1.28
 Consensus includes gb:AK025348.1 /DEF=Homo sapiens cDNA: FLJ21695 fls, clone COL09653. /FEA=mRNA /DB_XREF=gi:10437846 /UG=Hs.308803 Homo sapiens cDNA: AK025348 1.28
 diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein) NM_024104 1.28
 branched chain keto acid dehydrogenase E1, beta polypeptide (maple syrup urine disease) BC008466 1.28
 ribosomal protein L22 M55575 1.28
 transcription elongation factor 8 (SIII), polypeptide 1 (15kD, elongin C) NM_000983 1.28
 gb:Z25432.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405742 /FL=gb:Z25432. Z25432 1.28
 hypothetical protein, clone 24751 AF070530 1.28
 surfactant 5 NM_006752 1.28
 karyopherin (importin) beta 1 L38951 1.28
 BCR downstream signaling 1 NM_012103 1.28
 Consensus includes gb:Z88751 /DEF=Human DNA sequence from PAC 56089 on chromosome 1q24-q25. Contains profilin-like pseudogene. 60S ribosomal protein L4 pseud 298751. A4401703 1.28
 hypothetical protein FLJ22187 NM_017518 1.28
 Xq28. 2000bp sequence contig. ORF NM_024936 1.28
 hypothetical protein FLJ2024 AF177272 1.28
 UDP glycosyltransferase 2 family, polypeptide B28 BE384482 1.28
 SMARE protein NM_007213 1.28
 Consensus includes gb:AK025191.1 /DEF=Homo sapiens cDNA: FLJ21538 fls, clone COL08151. /FEA=mRNA /DB_XREF=gi:10437654 /UG=Hs.306783 Homo sapiens cDNA: AK025191 1.28
 JIM4 protein A1661768 1.28
 citron (Pro-interacting, serine/threonine kinase 21) BC005853 1.28
 hypothetical protein MGC15396 D87920 1.28
 solute carrier family 5 (sodium iodide symporter), member 5 AF164598 1.28
 CDC16 cell division cycle 16 homolog (S. cerevisiae) NM_000247 1.28
 MHC class I polypeptide-related sequence A AF164598 1.28
 CDC16 cell division cycle 16 homolog (S. cerevisiae) A1493119 1.28
 KIAA1198 protein AL385343 1.28
 solute carrier family 7 (carbonic amino acid transporter, y+ system), member 8 NM_003188 1.28
 mitogen-activated protein kinase kinase kinase 7 NM_002405 1.28
 manic fringe homolog (Drosophila) AB028973 1.28
 myelin transcription factor 1 BC004331 1.28
 hypothetical protein FLJ10814 AF217514 1.28
 hypothetical protein NM_018230 1.28
 lectin-cholesterol acyltransferase NM_000229 1.28
 three prime repair exonuclease 1 AJ243797 1.28
 carnitine palmitoyltransferase 1, muscle U62733 1.28
 hypothetical protein AL217472 1.28
 DNA-dependent protein kinase catalytic subunit-interacting protein 2 NM_006383 1.28
 Consensus includes gb:AC004144 /DEF=Homo sapiens chromosome 19, overlapping cosmids R28707 and R34001 /FEA=CDS /DB_XREF=gi:2887496 /UG=Hs.307350 Homo sapiens cDNA: AC004144 1.28
 integrin, beta 5 BE138575 1.28
 transmembrane trafficking protein NM_006827 1.28
 RA-regulated nuclear matrix-associated protein NM_016448 1.28
 desitin (actin depolymerizing factor) NM_006870 1.28
 protein phosphatase 1, regulatory (inhibitor) subunit 15A U63961 1.28
 DNKZP434F195 protein BE222901 1.28
 hypothetical protein FLJ10101 NM_024718 1.28
 Consensus includes gb:AF123462 /DEF=Homo sapiens BAC526N18 neurexin III gene, partial cds /FEA=CDS /DB_XREF=gi:4240565 /UG=Hs.247837 Homo sapiens BAC526N18 neurexin III gene, partial cds: AF123462 1.28
 hypothetical protein MGC3133 NM_031287 1.28
 hypothetical protein FLJ20718 AV584285 1.28
 melanin 2 NM_006554 1.28

RAN binding protein 2	D42063	128
adenosine A2a receptor	NM_000675	128
thyroid hormone receptor-associated protein, 95-kD subunit	AA314406	128
KIAA0947 protein	BC004902	128
son of sevenless homolog 1 (Drosophila)	L13857	128
cell division cycle 27	AL566824	128
CGI-40 protein	AA150165	128
elastase 2, neutrophil	NM_001972	128
RAP1, GTPase activating protein 1	NM_002865	128
KRAB-zinc finger protein SZF1-1	NM_016089	128
protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	BC006344	128
hypothetical protein FLJ23109	NM_024814	128
mitochondrial ribosomal protein L20	NM_017971	128
pancreatic lipase	NM_000936	128
bone morphogenetic protein receptor, type II (serine/threonine kinase)	U20165	128
hypothetical protein FLJ22189	NM_024085	128
ribosomal protein L23a	U43701	128
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	AJ224143	128
chromosome 1 open reading frame 16	BF509099	128
effector receptor, family 2, subfamily A, member 9 pseudogene	AA731709	128
ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	X69909	128
phospholipid transfer protein	NM_006227	128
splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)	NM_005066	128
hypothetical protein MGC5528	NM_024094	128
splicing factor, arginine/serine-rich 4	NM_005626	128
hypothetical protein FLJ20345	NM_017777	128
antigen p87 (melanoma associated) identified by monoclonal antibodies 133.2 and 96.5	NM_005929	128
hypothetical protein MGC2835	NM_024072	128
farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltransferase, geranyltransferase)	AL022163	128
protein phosphatase 1, regulatory subunit 7	A060331	128
CDC5 cell division cycle 5-like (S. pombe)	NM_001253	128
TFE3 target gene 1	BC002709	128
EH-domain containing 1	AY007161	128
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsinase, antitrypsin), member 2	NM_006220	128
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	NM_004725	128
muscle associated lymphoid tissue lymphoma translocation gene 1	NM_006785	127
RAB11A, member RAS oncogene family	AJ215102	127
RNA binding motif, single stranded interacting protein 2	D28483	127
gb:NM_016104.1 /DEF=Homo sapiens PTD013 protein (PTD013), mRNA, /FEA=mRNA /GEN=PTD013 /PROD=PTD013 /PROM=PTD013 /UG=Hs 279857 PTDc NM_016104	NM_016104	127
zinc finger protein 282	AA521508	127
mitochondrial ribosomal protein S18A	AB049952	127
eukaryotic translation initiation factor 4 gamma, 2	NM_001418	127
hypothetical protein FLJ23053	NM_022907	127
myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78)	NM_002462	127
baculoviral IAP repeat-containing 7 (livin)	NM_022161	127
sperm associated antigen 9	AK023512	127
ring finger protein 1	Z14000	127
hypothetical protein FLJ10587	NM_018149	127
cytosolic ovarian carcinoma antigen 1	AK000353	127
BCL2-associated athanogene 5	AA457021	127
artuin alien mating type information regulation 2 homolog 3 (S. cerevisiae)	AW298092	127
KIAA0776 protein	AK023092	127
DNF2P436D2223 protein	BE567813	127
ATPase, Cu++ transporting, alpha polypeptide (Mankes syndrome)	NM_006844	127
Irb (bacterial scitadicate synthase)-like	AB041926	127
Mishapen/NIK-related kinase	AI373539	127
HIV TAT specific factor 1	AK023421	127
SEC7 homolog	AW873594	127
hypothetical protein	AW873594	127
Consensus includes gb:AW873564 /FEA=EST /DB_XREF=gi:8007617 /DB_XREF=eat:h62c10.x1 /CLONE=IMAGE:3041970 /UG=Hs 155485 huntingtin interacting protein 2	NM_004376	127
COX15 homolog, cytochrome c oxidase assembly protein (yeast)	NM_002056	127
glutamine-fructose-6-phosphate transaminase 1	NM_025097	127
hypothetical protein FLJ21106		127

inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta
membrane-bound transcription factor protease, site 2
homoeo box (H6 family) 1
splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)
serine palmitoyltransferase, long chain base subunit 1
hypothetical protein
KIAA1288 protein
KIAA1698 protein
hypothetical protein FLJ10143
RAB8B, member RAS oncogene family
KIAA0212 gene product
ubiquitin specific protease 8
phosphatase and leucine homolog (mutated in multiple advanced cancers 1)
amino acid transporter system A1
aplatxin
plekrophilin 3
exportin 1 (CRM1 homolog, yeast)
hypothetical protein FLJ20986
ubiquitin 1
hypothetical protein FLJ10659
alpha disintegrin and metalloproteinase domain 23
heterogeneous nuclear ribonucleoprotein A0
testis enhanced gene transcript (BAX inhibitor 1)
PC2 (positive cofactor 2, multiprotein complex) glutamine/O-rich-associated protein
GTP binding protein 1
splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3
gb:NM_015803.1 /DEF=Homo sapiens transcriptional intermediary factor 1 (TIF1), mRNA, /FEA=mRNA /GEN=TIF1 /PROD=transcriptional intermediary factor 1 alpha /DB_XREF=NM_015803
synapobrevin-like 1
5-oxoprolinase (ATP-hydrolysing)
actin related protein 2/3 complex, subunit 2 (34 kD)
KIAA0423 protein
albumin
proteasome (prosome, macropain) subunit, beta type, 1
Consensus includes gb:AW970944 /FEA=EST /DB_XREF=gi:8160789 /DB_XREF=est:EST333027 /UG=Hs.291839 ESTs
KIAA0989 protein
KIAA0304 gene product
KIAA0392 protein
Consensus includes gb:A287546 /FEA=EST /DB_XREF=gi:3886713 /DB_XREF=est:est:aq8005.x1 /CLONE=IMAGE:2038209 /UG=Hs.90798 Human clone 23695 mRNA sequen
secretory carrier membrane protein 1
aspartate beta-hydroxylase
hypothetical protein FLJ22195
mannose-6-phosphate utilization defect 1
annexin A7
p100 protein interacting protein
synaptobrevin V
ribosomal protein L22
gila maturation factor, beta
T54 protein
hypothetical protein FLJ20729
heat shock 90kD protein 1, alpha
BAI1-associated protein 3
protein kinase, AMP-activated, beta 1 non-catalytic subunit
interferon, gamma-inducible protein 30
culin 2
gb:NM_020412.1 /DEF=Homo sapiens CHMP1.5 protein (CHMP1.5), mRNA, /FEA=mRNA /GEN=CHMP1.5 /PROD=CHMP1.5 protein /DB_XREF=gi:9866900 /UG=Hs.42733 (NM_020412
major histocompatibility complex, class I, F
Consensus includes gb:AA931266 /FEA=EST /DB_XREF=gi:3085652 /DB_XREF=est:om8105.s1 /CLONE=IMAGE:1554273 /UG=Hs.315478 Homo sapiens, Similar to perlecan
hypothetical protein MGC11138
insulin promoter factor 1
insulin promoter factor 1, homeodomain transcription factor
joined to JAZF1
protein (peptidyl-prolyl isomerase) NIMA-interacting 1
Consensus includes gb:AK024456.1 /DEF=Homo sapiens mRNA for FLJ00048 protein, partial cds, /FEA=mRNA /GEN=FLJ00048 /PROD=FLJ00048 protein /DB_XREF=gi:104 AK024456

hypothetical protein FLJ10008 1.26
 small GTP-binding protein 1.26
 RNA 3'-terminal phosphate cyclase 1.26
 H1 histone family, member 2 1.26
 Consensus includes gb-BG178274 /FEA=EST /DB_XREF=gi:12684977 /DB_XREF=est:502330318f1 /CLONE=IMAGE:4431676 /UG=Hs.278895 postmeiotic segregation Invert BG178274 1.26
 Consensus includes gb-AA527409 /FEA=EST /DB_XREF=gi:2269568 /DB_XREF=est:ng4107.s1 /CLONE=IMAGE:937381 /UG=Hs.301667 Homo sapiens mRNA; cDNA DKF:AA527499 1.26
 heterogeneous nuclear ribonucleoprotein U (scarfold attachment factor A) 1.26
 translocase of inner mitochondrial membrane 22 homolog (yeast) 1.26
 Consensus includes gb-AL121777 /DEF=Human DNA sequence from clone RPS-1057D4 on chromosome 20 Contains a Spermidine synthase (SPDSY) pseudogene, a CpG is AL121777 1.26
 cullin 4A 1.26
 RNA (guanine-7-) methyltransferase 1.26
 arrestin, beta 1 1.26
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 1.26
 hypothetical protein DC30 1.26
 poly(A) binding protein, cytoplasmic 3 1.26
 hypothetical protein R33729.1 1.26
 GTPase regulator associated with the focal adhesion kinase pp125 1.26
 N-acetylglucosamine-phosphate mutase 1.26
 cutaneous T-cell lymphoma-associated tumor antigen se20-4, differentially expressed nucleolar TGF-beta 1 target protein (DENT1) 1.26
 protocadherin beta 1 1.26
 hypothetical protein FLJ22405 1.26
 Consensus includes gb-AWA09974 /FEA=EST /DB_XREF=gi:6935515 /DB_XREF=est:fn03d12.y1 /CLONE=IMAGE:2961094 /UG=Hs.332750 Homo sapiens cDNA FLJ10100 1.26
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a-like 1 1.26
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 1.26
 uncharacterized bone marrow protein BM029 1.26
 zinc finger protein 197 1.26
 KIAA1491 protein 1.26
 cytochrome c oxidase subunit VIc 1.26
 dynein light chain-A 1.26
 Consensus includes gb-AA310709 /FEA=EST /DB_XREF=gi:1963058 /DB_XREF=est:EST181521 /UG=Hs.249187 Homo sapiens BAC clone RP11-121A8 from 7p14-p13 1.26
 EH-domain containing 1 1.26
 apolipoprotein E 1.26
 gqgi SNAP receptor complex member 2 1.26
 dual specificity phosphatase 12 1.26
 laminin receptor 1 (87kD, ribosomal protein SA) 1.26
 T4 translation elongation factor, mitochondrial 1.26
 hypothetical protein FLJ13909 1.26
 NAD kinase 1.26
 chromosome 22 open reading frame 2 1.26
 hypothetical protein FLJ10709 1.26
 karyopherin (importin) beta 1 1.26
 actin cross-linking factor 1.26
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) 1.26
 diaphenous homolog 1 (Drosophila) 1.26
 MAOS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C) 1.26
 calcium channel, voltage-dependent, gamma subunit 2 1.26
 Cbort10 protein 1.26
 nuclear receptor subfamily 8, group A, member 1 1.26
 flamin A, alpha (actin binding protein 280) 1.26
 hypothetical protein FLJ21272 1.26
 zinc finger RNA binding protein 1.26
 KIAA1411 protein 1.26
 hypothetical protein FLJ11749 1.26
 microsomal glutathione S-transferase 2 1.26
 DKF ZP547E2110 protein 1.26
 hypothetical protein FLJ10704 1.26
 ganglioside-induced differentiation-associated protein 1 1.26
 polycyochrome c synthase (cytochrome c heme-lyase) 1.26
 folypolyglutamate synthase 1.26
 Williams-Beuren syndrome chromosome region 1 1.26
 hypothetical protein FLJ13439 1.26
 Consensus includes gb-AI078186 /FEA=EST /DB_XREF=gi:3405364 /DB_XREF=est:oz01g01.x1 /CLONE=IMAGE:1674096 /UG=Hs.284275 Homo sapiens PAK2 mRNA, com 1.26
 beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase 1) 1.26

RAS p21 protein activator 2	NM_005506	1.26
keratin alpha 1 (Importin alpha 5)	BC02374	1.26
germ cell specific Y-box binding protein	NM_015982	1.26
nuclear cap binding protein subunit 1, 80kD	BC00193	1.26
eukaryotic translation initiation factor 3, subunit 6 (48kD)	BC000734	1.26
BTG family, member 2	BC339064	1.26
CD4-141 protein	NM_016072	1.26
secretory carrier membrane protein 1	AV745949	1.25
nuclear factor I/B	U70862	1.25
NEL sepin-like fusion	NM_006640	1.25
homo box B5	A052747	1.25
Consensus includes gb:AI017564 /FEA=EST /DB_XREF=gi:3231900 /DB_XREF=est:cu23e03.x1 /CLONE=IMAGE:1627132 /UG=Hs.95631 Human normal keratinocyte mRNA	AI017564	1.25
patched homolog (Drosophila)	BC054916	1.25
Consensus includes gb:AW519168 /FEA=EST /DB_XREF=gi:7157250 /DB_XREF=est:ha52a08.x1 /CLONE=IMAGE:2877302 /UG=Hs.163271 Human alpha-1 Ig germline C- α	AW519168	1.25
KIAA0378 protein	AB002374	1.25
cyclchrome c oxidase subunit Va pseudogene 1	AF042162	1.25
golgi SNAP receptor complex member 2	BE895285	1.25
hypothetical protein FLJ20288	NM_024668	1.25
Consensus includes gb:AW452796 /FEA=EST /DB_XREF=gi:5993572 /DB_XREF=est:UH-H-B13-alk-4-06-0-UI.s1 /CLONE=IMAGE:3069059 /UG=Hs.120769 Homo sapiens cDN	AW452796	1.25
hsp70-interacting protein	NM_012287	1.25
KIAA0033 protein	AU153138	1.25
serine protease inhibitor, Kazal type 1	NM_003122	1.25
CDC23 (cell division cycle 23, yeast, homolog)	NM_004661	1.25
KIAA0728 protein	AF798790	1.25
BRCA1 associated protein	AL042733	1.25
Consensus includes gb:AV691323 /FEA=EST /DB_XREF=gi:10293186 /DB_XREF=est:AV691323 /CLONE=GKCEWF11 /UG=Hs.2056 UDP glycosyltransferase 1 family, poly	AV691323	1.25
KIAA0923 protein	AW136988	1.25
putative nucleic acid binding protein RY-1	X76302	1.25
hippocampin-like 1	BE917568	1.25
KIAA0350 protein	AA731693	1.25
hypothetical protein FLJ13912	AI421812	1.25
glyoxalase 1	NM_006708	1.25
erythropoietin	AF053356	1.25
casein kinase 1, alpha 1	AF118911	1.25
fibronectin 1	AK026737	1.25
Hsp90-associated relative of Cdc37	NM_017913	1.25
zinc finger protein 134 (clone pRZ-15)	NM_003435	1.25
PR domain containing 2, with ZNF domain	AV724783	1.25
argininosuccinate lyase	NM_000048	1.25
KIAA0828 protein	AB020635	1.25
platelet derived growth factor C	NM_016205	1.25
golgi autoantigen, golgin subfamily a, 5	NM_005113	1.25
RPB5-mediating protein	AW514900	1.25
granulin	AK023348	1.25
endothelin receptor type B	M74921	1.25
HEMLK homolog 7b	W93807	1.25
aminoacidipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase	AF151057	1.25
hypothetical protein FLJ14872	AA535065	1.25
solute carrier family 18 (monocarboxylic acid transporters), member 3	AA853175	1.25
cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)	BC000059	1.25
WNT1 inducible signaling pathway protein 2	NM_003881	1.25
small inducible cytokine subfamily B (Cys-X-Cys), member 11	AF002985	1.25
chromogranin A (parathyroid secretory protein 1)	NM_001275	1.25
CLIP-associated protein 1	AA176798	1.25
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs	AI828531	1.25
Consensus includes gb:AW449390 /FEA=EST /DB_XREF=gi:6990166 /DB_XREF=est:UH-H-B13-alk-4-06-0-UI.s1 /CLONE=IMAGE:2734650 /UG=Hs.257150 ESTs, Moderately	AW449390	1.25
signal transducer and activator of transcription 6, interleukin-4 induced	BC004973	1.25
sumo1 homolog 1 (Drosophila)	NM_014676	1.25
CDC-like kinase 3	NM_003992	1.25
polycystic kidney disease 1 (autosomal dominant)	NM_000296	1.25
TLA1 cytosolic granule-associated RNA binding protein-like 1	NM_003252	1.25
isoprenylcysteine carboxyl methyltransferase	NM_012405	1.25
ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)	NM_020469	1.25

A20-binding inhibitor of NF-kappaB activation-2
 Consensus includes gb:269637.1 /DEF=H.sapiens mRNA sequences (16p11.2). /FEA=mRNA /DB_XREF=gi:29573 /UG=Hs.307208 H.sapiens mRNA sequence (16p11.2)
 uncharacterized hematopoietic stem/progenitor cells protein MDS033
 Rho guanine exchange factor (GEF) 11
 Consensus includes gb:A1852772 /FEA=EST /DB_XREF=gi:5745082 /DB_XREF=est:ws6005.x1 /CLONE=IMAGE:2491113 /UG=Hs.300865 Homo sapiens clone BCSynL38
 Consensus includes gb:A1852772 /FEA=EST /DB_XREF=gi:5745082 /DB_XREF=est:ws6005.x1 /CLONE=IMAGE:2491113 /UG=Hs.300865 Homo sapiens clone BCSynL38
 transmembrane 4 superfamily member 6
 hypothetical protein FLJ20030
 growth arrest and DNA-damage-inducible, alpha
 HSPC133 protein
 hypothetical protein FLJ22347
 luciferase 2 (secretor status included)
 diacylglycerol kinase, beta
 uncharacterized hypothalamus protein HT007
 protein phosphatase 3 (formy 2B), catalytic subunit, beta isoform (calcineurin A beta)
 BCL2adenovirus E1B 19kD interacting protein 1
 forhead box H1
 ATP-binding cassette, sub-family G (WHITE), member 1
 DKFZP568B1621 protein
 HSPC134 protein
 carboxypeptidase D
 hypothetical protein FLJ11273
 x 005 protein
 chromodomain helicase DNA binding protein 4
 hypothetical protein FLJ14390
 myeloid lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10
 spinlin
 zinc finger protein 108
 branched chain alpha-ketoadid dehydrogenase kinase
 cell division cycle 2-like 1 (PTSLRE proteins)
 hypothetical protein FLJ13910
 phospholipase C, gamma 2 (phosphatidylinositol-specific)
 carboxypeptidase Z
 hypothetical protein BC022942
 Consensus includes gb:AL390153.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762G113 (from clone DKFZp762G113). /FEA=mRNA /DB_XREF=gi:9368841 /UG=Hs.208339
 hypothetical protein FLJ20561
 baculoviral AP repeat-containing 1
 candidate mediator of the p53-dependent G2 arrest
 putative DNA binding protein
 MCM6 minichromosome maintenance deficient 6 (MISS homolog, S. pombe) (S. cerevisiae)
 ARP1 actin-related protein 1 homolog B, centractin beta (yeast)
 nasopharyngeal epithelium specific protein 1
 KIAA0354 protein
 heterogenous nuclear ribonucleoprotein H1 (H)
 hypothetical protein FLJ20548
 retinoic acid induced 16
 beta-2-microglobulin
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 11
 eductin 3 (gamma)
 polymerase (DNA directed), mu
 KIAA1098 protein
 dynamin 1
 copin VI (neuronal)
 hypothetical protein FLJ10208
 hypothetical protein MGC3282
 hypothetical protein FLJ10743
 suppressor of potassium transport defect 3
 tumor differentially expressed 1
 Consensus includes gb:S79287.1 /DEF=CD4 receptor (exons 1 and 2) human, T-lymphocyte, mRNA, 3428 nt. /FEA=mRNA /GEN=CD4 receptor /PROD=CD4 receptor /DB_XS_S79287
 Splicing factor, arginine/leucine-rich, 46kD
 hypothetical protein FLJ22035
 pancretic polypeptide
 Consensus includes gb:AW12802.1 /FEA=EST /DB_XREF=gi:8116965 /DB_XREF=est:xe93h03.x1 /CLONE=IMAGE:2615477 /UG=Hs.243537 ESTs, Highly similar to AC00504
 JMS protein
 BC000464

KIAA0993 protein
 hypothetical protein FLJ10379
 tyrosinase (H. influenzae) homolog
 tubulin beta-5
 hypothetical gene MGC9753
 cytosine conjugate-beta lyase, cytoplasmic (glutamine transaminase K, kynurenine aminotransferase)
 signal transducing adaptor molecule (SH3 domain and ITAM motif) 1
 ribosomal protein S8
 BCL2-associated X protein
 COX15 homolog, cytochrome c oxidase assembly protein (yeast)
 hypothetical protein HSPC177
 putative-rich element binding protein A
 centrosomal protein 2
 paired box gene 2
 hypothetical protein DJ12208.2
 Cas-B1-M (murine) ectropic retroviral transforming sequence c
 Cas2-promoted Ras inactivator
 Consensus includes gb:AK000834.1 /DEF=Homo sapiens cDNA FLJ20827 flt. clone ADKA03543. /FEA=mrna /DB_XREF=gi:7021152 /UG=Hs.306409 Homo sapiens cDNA
 soluble carrier family 7 (cationic amino acid transporter, y+ system), member 8
 fatty acid desaturase 1
 hypothetical protein FLJ20327
 Consensus includes gb:AL035588 /DEF=Human DNA sequence from clone 696P19 on chromosome 6p12.3-21.2. Contains the gene for TFEB, an NPM1 (Nucleophosmin, Nur
 hypothetical protein FLJ10385
 homeo box B5
 LRP16 protein
 KIAA0102 gene product
 FYT0 domain-containing ion transport regulator 1 (phospholemman)
 Consensus includes gb:AL583123 /FEA=EST /DB_XREF=gi:12951784 /DB_XREF=est:AL583123 /CLONE=CSODL009YN08 (3 prime) /UG=Hs.294005 Homo sapiens, clone
 hypothetical protein FLJ13220
 topoisomerase (DNA) II alpha (170kD)
 protocadherin gamma subfamily A, 10
 polymerase (DNA directed), epsilon 2
 autophagic translation initiation factor 3, subunit 3 (gamma, 40kD)
 microtubule-associated protein 4
 zinc finger protein 145 (Kruppel-like, expressed in promyelocytic leukemia)
 zinc finger protein 14
 CQI-42 protein
 annexin A2 pseudogene 2
 Consensus includes gb:BF448531 /FEA=EST /DB_XREF=gi:11514556 /DB_XREF=est:17n72e02.x1 /CLONE=IMAGE:3570146 /UG=Hs.248979 ESTs, Moderately similar to ALI
 ribosomal protein L22
 hypothetical protein FLJ11159
 ARF3 actin-related protein 3 homolog (yeast)
 KIAA0116 protein
 Huntingtin interacting protein K
 DKFZP434B103 protein
 KIAA0155 gene product
 SFY (sex determining region Y)-box 11
 diacylglycerol O-acyltransferase homolog 1 (mouse)
 adenomatous polyposis coli
 ring finger protein 24
 ring finger protein 24
 cylindromatosis (turban tumor syndrome)
 chromosome 11 open reading frame 13
 thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)
 choline kinase
 hypothetical protein FLJ20258
 Consensus includes gb:A0807507 /FEA=EST /DB_XREF=gi:5394073 /DB_XREF=est:W448e10.x1 /CLONE=IMAGE:2358858 /UG=Hs.126496 Homo sapiens chromosome 19, c
 TRAM-like protein
 protein phosphatase 1, catalytic subunit, gamma isoform
 nucleoporin 50KD
 zinc finger protein, subfamily 1A, 1 (Ikaros)
 KIAA0552 gene product
 synaptic vesicle protein 28 homolog

synovial sarcoma, X breakpoint1 2	BC002818	1.17
fibroblast growth factor 8	NM_020986	1.17
opoid binding protein/cell adhesion molecule-like	NM_002545	1.17
G protein-coupled receptor 44	AF118265	1.17
Fzr1 protein	AB013463	1.17
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 2	NM_002070	1.17
outstative 28 kDa protein	NM_020143	1.17
like-glycosyltransferase	AB011161	1.17
lanthipeptin (importin) beta 2	NM_002270	1.17
cell division cycle 42 (GTP binding protein, 25kD)	R37664	1.17
SCO cytochrome oxidase deficient homolog 2 (yeast)	NM_005138	1.17
jagged 2	AF029778	1.17
v-rat-1 murine leukemia viral oncogene homolog 1	X09409	1.17
frizzled homolog 5 (Drosophila)	NM_003468	1.17
Consensus includes gb:H24302 /FEA=EST /DB_XREF=gi:892997 /DB_XREF=est:ym51e07.s1 /CLONE=IMAGE:51738 /UG=Hs.23127 ESTs	H24302	1.17
hypothetical protein MGC4276 similar to CG8188	BC002675	1.17
Consensus includes gb:A1188739 /FEA=EST /DB_XREF=gi:3737377 /DB_XREF=est:q78c01.x1 /CLONE=IMAGE:1745184 /UG=Hs.326416 Homo sapiens mRNA; cDNA DKF	A1188739	1.17
MYC-associated zinc finger protein (putative-binding transcription factor)	A471665	1.17
hypothetical protein FLJ20657	NM_017921	1.17
splicing factor, arginine/serine-rich 2, interacting protein	A1954932	1.17
hypothetical protein FLJ12448	NM_022895	1.17
G antigen, family B, 1 (prostate associated)	NM_003785	1.17
GDP dissociation inhibitor 1	NM_001493	1.17
Consensus includes gb:AL137784 /DEF=Human DNA sequence from clone RP1-189J3 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains part of the	NM_003785	1.17
Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704480 /UG=Hs.295689 H.sapiens clathrin light chain a gene	NM_003785	1.17
Ras-GTPase activating protein SH3 domain-binding protein 2	AB014560	1.17
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, E116, E130, EL32 and G344)	BF983379	1.17
hypothetical protein, similar to (U06944) PRAJA1	AL530462	1.17
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2	NM_002575	1.17
melanoma antigen, family D, 1	AF217963	1.17
endosome-associated FYVE-domain protein	NM_014723	1.17
muclin and cadherin-like	NM_021924	1.17
ROD1 regulator of differentiation 1 (S. pombe)	AW190873	1.17
KIAA1696 protein	NM_016621	1.17
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease)	AF077973	1.17
G protein-coupled receptor 51	AF095723	1.17
alko-keto reductase family 1, member C3 (3-alpha hydroxysteroid dehydrogenase, type II)	AB018580	1.17
leucocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	U82276	1.17
origin recognition complex, subunit 4-like (yeast)	NM_002552	1.17
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14)	NM_002490	1.17
mitofusin 1	NM_017827	1.17
hypothetical protein FLJ10545	NM_018132	1.17
integral membrane protein 1	NM_002219	1.17
G protein-coupled receptor 56	NM_005682	1.17
JAK binding protein	AA877218	1.17
stem cell growth factor: lymphocyte secreted C-type lectin	NM_002975	1.17
heterogeneous nuclear ribonucleoprotein A3	NM_005758	1.17
G protein-coupled receptor 4	U35399	1.17
mitochondrial ribosomal protein S31	NM_005830	1.17
Consensus includes gb:AF189251.1 /DEF=Homo sapiens cytomegalovirus partial fusion receptor mRNA, partial cds. /FEA=mRNA /PROD=cytomegalovirus partial fusion recar	AF189251	1.17
SAC2 suppressor of actin mutations 2-like (yeast)	AL390171	1.17
KIAA0751 gene product	AF007156	1.17
protein phosphatase 2 (formerly 2A), regulatory subunit B' (PR 72), alpha isoform and (PR 130), beta isoform	NM_002718	1.17
polymerase (RNA) II (DNA directed) polypeptide K (7.0kD)	AL558030	1.17
S-adenosylhomocysteine hydrolase-like 1	NM_006621	1.17
excision repair cross-complementing rodent repair deficiency, complementation group 3 (xeroderma pigmentosum group B complementing)	NM_000122	1.17
dihydropyrimidine branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease)	M27093	1.17
calcium binding atopy-related autoantigen 1	AK022697	1.17
phenylalanine-tyrosine synthetase	NM_006567	1.17
hypothetical protein MGC4808	NM_024516	1.17
SRB7 suppressor of RNA polymerase B homolog (yeast)	U46837	1.17
3-hydroxyethyl-3-methylglutaryl-Coenzyme A lyase (hydroxymethylglutaryl-CoA lyase)	NM_000191	1.17
gastric inhibitory polypeptide	NM_004123	1.17

hypothetical SBB103 protein	AL583171	1.17
cardio-lipofuscinosis, neuronal 5	A911687	1.17
hypothetical protein MGC10765	NM_024345	1.17
Rho guanine nucleotide exchange factor (GEF) 5	NM_005435	1.17
X-ray repair complementing defective repair in Chinese hamster cells 3	AK022829	1.17
KIAA1155 protein	AB032981	1.17
similar to yeast BET3 (S. cerevisiae)	NM_014408	1.17
mitogen-activated protein kinase kinase 5	NM_002757	1.17
DMP27721M231 protein	AL117480	1.17
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AI684141	1.17
similar to yeast Upt3, variant A	NM_023011	1.17
heart shock 70kD protein 4	AB023420	1.17
cannabinoid receptor 1 (brain)	NM_016083	1.17
gpJF308302.1 IDEF-Homo sapiens serologically defined breast cancer antigen NY-BR-86 mRNA, complete cds. /FEA=mRNA PRO-ol-serologically defined breast cancer ant	AF308302	1.17
KIAA1058 protein	AY450751	1.17
protease, cysteine, 1 (legumain)	D55896	1.17
type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	AF106037	1.17
histone deacetylase 5	NM_005474	1.17
ring finger protein 27	NM_030912	1.17
tumor endothelial marker 1 precursor	NM_020404	1.17
collagen, type IV, alpha 3 (Goodpasture antigen) binding protein	NM_005713	1.17
mitogen-activated protein kinase-activated protein kinase 2	BC000351	1.17
phosphate cytidylyltransferase 2, ethanolamine	AB035745	1.17
Down syndrome critical region gene 5	NM_005781	1.17
activated p21cdc42Hs kinase	NM_007284	1.17
protein tyrosine kinase 9-like (A6-related protein)	NM_022776	1.16
hypothetical protein FLJ13164	NM_021082	1.16
solute carrier family 15 (H+/peptide transporter), member 2	BE350145	1.16
regulator of nonsense transcripts 2	NM_015542	1.16
DKF ZP564A043 protein	AL036344	1.16
mucin 6, tracheobronchial	U14383	1.16
zinc finger protein 148 (pH2.52)	L04282	1.16
splicing factor, arginine/serine-rich 9	NM_003769	1.16
numa homolog (Drosophila)	NM_003744	1.16
PBX/Notch1 1 homeobox 1	NM_004571	1.16
peroxisome receptor 1	AW468717	1.16
PTK6 protein tyrosine kinase 6	NM_005975	1.16
sorcin	L12387	1.16
adaptor-related protein complex 3, beta 1 subunit	AW058575	1.16
hypothetical protein FLJ10781	NM_018215	1.16
protein phosphatase 8, catalytic subunit	NM_018294	1.16
shvelar soft part sarcoma chromosome region, candidate 1	NM_024083	1.16
zinc finger protein 95 homolog (mouse)	NM_014589	1.16
XIAP associated factor-1	NM_017523	1.16
cytochrome P450, subfamily XXIV (vitamin D 24-hydroxylase)	AI432196	1.16
general transcription factor II, 1	NM_000782	1.16
zinc finger protein, subfamily 1A, 3 (Atf10s)	AF035737	1.16
glycine cleavage system protein H (aminomethyl carrier)	NM_012481	1.16
transforming growth factor, beta 2	NM_006311	1.16
Conserved gene telomeric to alpha globin cluster	AW237404	1.16
glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID)	NM_003238	1.16
TNF receptor-associated factor 2	BC004185	1.16
hypothetical protein FLJ22893	NM_002076	1.16
Kruppel-like factor 15	NM_021138	1.16
pleomorphic adenoma gene-like 1	NM_022750	1.16
platelet-activating factor acetylhydrolase, isoform lb, alpha subunit (45kD)	NM_014079	1.16
glucosamine-6-phosphate isomerase	NM_006718	1.16
KIAA0852 protein	L13386	1.16
zinc finger protein 140 (clone pHZ-39)	NM_005471	1.16
KIAA0152 gene product	AC004542	1.16
	NM_003440	1.16
	BC000371	1.16

protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform
homo box D3
DnaJ (Hsp40) homolog, subfamily B, member 6
Kruppel-like factor 3 (Intestinal)
fragile X mental retardation 1
cyclin L, anla-6a
karyopherin (importin) beta 2
UDP-N-acetylglucosamine pyrophosphorylase 1
cell division cycle 25b
FtsJ homolog 2 (E. coli)
non-metastatic cells 3, protein expressed in
progesterone receptor membrane component 1
Mouse Mammary Tumor Virus Receptor homolog
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2
ubiquitin specific protease 15
chloride intracellular channel 1
KIAA0871 gene product
hypothetical protein MGC955
cell cycle progression 2 protein
olfactory receptor, family 2, subfamily F, member 1
epimorphin
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform
gb:NM_016811.1/DEF=Homo sapiens hypothetical protein PRO1966 (PRO1966), mRNA, /FEA=mRNA /GEN=PRO1966 /PROD=hypothetical protein PRO1966 /DB_XREF=gi:10203952 /DB_XREF=est:601471903F1 /CLONE=IMAGE:2804508 /UG=Hs.102178 syntaxin 16 /FL=gb:AF00893-BC762754
cyclin M3
KIAA0560 gene product
hypothetical protein FLJ11786
signal peptidase complex (18kD)
potassium large conductance calcium-activated channel, subfamily M beta member 3
sodium channel, nonvoltage-gated 1, gamma
HIV-1 Tat interactive protein, 60 kDa
sarcosine dehydrogenase
tumor necrosis factor receptor superfamily, member 6b, decoy
SPTF-associated factor 65 gamma
platelet-derived growth factor alpha polypeptide
breast cancer 1, early onset
karyopherin (importin) beta 1
microtubule-associated protein, RP/EB family, member 2
tumor protein D52
hypothetical protein FLJ13611
KIAA0157 protein
adaptor-related protein complex 1, sigma 1 subunit
Consensus includes gb:BE762754 /FEA=EST /DB_XREF=gi:10203952 /DB_XREF=est:601471903F1 /CLONE=IMAGE:3874921 /UG=Hs.102178 syntaxin 16 /FL=gb:AF00893-BC762754
ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)
Isovaleryl Coenzyme A dehydrogenase
zinc finger protein 215
adducin 3 (gamma)
kinesin-like 2
neuropeptide Y receptor Y1
immunoglobulin (CD79A) binding protein 1
neuraxin 1
XPM12 protein
signaling lymphocytic activation molecule
acyl-Coenzyme A oxidase 2, branched chain
arachidonate 5-lipoxygenase
transport-secretion protein 2.2
ethanolamine kinase
Consensus includes gb:AW438464 /FEA=EST /DB_XREF=gi:6973770 /DB_XREF=est:u43g07.x1 /CLONE=IMAGE:2804508 /UG=Hs.288760 Homo sapiens cDNA FLJ12327 AW438464
gb:M5258.1 /DEF=Homo sapiens immunoglobulin kappa-chain VK-1 (IgK) mRNA, complete cds. /FEA=mRNA /GEN=IgK /PROD=immunoglobulin kappa-chain VK-1 /DB_XRI M5258
arachidonate 12-lipoxygenase pseudogene 2
complement component 4B
cyclin-dependent kinase (CDC2-like) 10
hypothetical protein FLJ11101
glucocorticoid modulatory element binding protein 2

NIMA (never in mitosis gene a)-related kinase 1	AI938517	1.16
trypsin motif-containing 10	X90539	1.16
trypsin motif-containing 2	AW071795	1.16
double homeobox, 2	NM_012147	1.16
CD209 antigen	AF260866	1.16
phospholipase A2, group IVB (cytosolic)	NM_005090	1.16
hypothetical protein FLJ21877	NM_022065	1.16
phosphatidylcholine transfer protein	NM_021213	1.16
KIAA0252 protein	D87440	1.16
Consensus includes gb:AK021738.1 /DEF=Homo sapiens cDNA FLJ11876 fis, clone HEMBA1004752, highly similar to Homo sapiens mRNA for LAX-4p. /FEA=mRNA /DB_XF AK021738		1.16
Consensus includes gb:Y15724 /DEF=Homo sapiens SERCA3 gene, exons 1-7 (and joined CDS) /FEA=mRNA /DB_XREF=gi:3021395 /UG=Hs.5541 ATPase, Ca++ transport Y15724		1.16
uncharacterized bone marrow protein BM046	NM_018460	1.16
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	BE767213	1.16
protein kinase C, eta	NM_006255	1.16
inter-alpha (globulin) inhibitor, H2 polypeptide	NM_002216	1.16
KIAA1719 protein	AF052177	1.16
hypothetical protein FLJ23338	NM_024658	1.16
heterogeneous nuclear ribonucleoprotein H1 (H)	NM_024658	1.16
ribulose-5-phosphate-3-epimerase	BF983406	1.16
hypothetical protein DKFZp434L0117	BE984473	1.16
glyoxyltransferase	NM_022778	1.16
microtubular-associated protein 4	NM_031302	1.16
Consensus includes gb:AF007131.1 /DEF=Homo sapiens clone 23963 mRNA sequence /FEA=mRNA /DB_XREF=gi:2852805 /UG=Hs.48483 Homo sapiens clone 23963 mR	R72286	1.16
mitogen-activated protein kinase kinase kinase 1	AA744529	1.16
5-hydroxytryptamine (serotonin) receptor 5A	NM_024012	1.16
CD72 antigen	AF283777	1.16
dual specificity phosphatase 5	U16996	1.16
Consensus includes gb:AL110206.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N2022 (from clone DKFZp586N2022) /FEA=mRNA /DB_XREF=gi:5817125 /UG=Hs.30633	AL110206	1.16
fibroblast growth factor 12B	NM_004113	1.16
calreticulin	BE251303	1.16
ribosomal protein S6 kinase, 90kD, polypeptide 1	NM_002853	1.16
LIM domain kinase 2	NM_005569	1.16
Consensus includes gb:Y08772 /DEF=H. sapiens PABII pseudogene /FEA=CDS /DB_XREF=gi:1770485 /UG=Hs.247983 H.sapiens PABII pseudogene	Y08772	1.16
fer-1-like 3, myoferlin (C. elegans)	AF207990	1.16
amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AV681579	1.16
related RAS viral (v-ras) oncogene homolog, 2	A431843	1.16
Consensus includes gb:AK026867.1 /DEF=Homo sapiens cDNA: FLJ23214 fis, clone ADSU01585. /FEA=mRNA /DB_XREF=gi:10439827 /UG=Hs.306891 Homo sapiens cDNA: AK026867	AK026867	1.16
inositol 1,4,5-trisphosphate receptor, type 3	NM_002224	1.16
myelin protein zero-like 1	AF095727	1.16
potassium voltage-gated channel, shaker-related subfamily, beta member 2	AF044253	1.16
COPG (constitutive photomorphogenic, Arabidopsis, homolog) subunit 4	NM_016129	1.16
TXK tyrosine kinase	NM_003328	1.16
epidermal growth factor receptor pathway substrate 15	BF213575	1.16
hypothetical protein R31240.1	A1435954	1.16
DEADH (Asp-Glu-Ala-Asp-His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae)	NM_004399	1.16
phosphatidylcholine-specific phospholipase D1	L11702	1.16
immunoglobulin heavy constant gamma 3 (G3m marker)	NM_025072	1.16
hypothetical protein FLJ14038	X95748	1.16
potassium channel, subfamily K, member 10 (TREK-2)	A1765051	1.16
PC4 and SFRS1 interacting protein 2	NM_021161	1.16
similar to KIAA0010 gene product (H. sapiens).	AF063020	1.16
hypothetical protein FLJ12517	AL096740	1.16
hypothetical protein FLJ10283	NM_023007	1.15
caspase 3, apoptosis-related cysteine protease	NM_018046	1.15
matrix metalloproteinase 19	NM_004346	1.15
KIAA1054 protein	U38321	1.15
calmodulin 2 (phosphorylase kinase, delta)	AB028977	1.15
protein tyrosine kinase 2 beta	NM_001743	1.15
blast cell autoantigen 1 (BRX)	U43522	1.15
enigma (LIM domain protein)	W67992	1.15
steroid regulatory element binding transcription factor 1	AA086229	1.15
multifunctional polypeptide similar to SAICAR synthetase and AIR carboxylase	NM_004176	1.15
	AA026552	1.15

kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
 glutathione peroxidase 4 (phospholipid hydroperoxidase)
 chromatin assembly factor 1, subunit B (p60)
 Custer Ind. AW025284: wu85h.10.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:990806 /clone_end=3 /gb=AW025284 /gi=5878814 /ug=Hs.237946 /len=454
 hemochromatosis
 nuclear receptor subfamily 6, group A, member 1
 KIAA0878 protein
 vesicle-associated membrane protein 2 (synaptobrevin 2)
 hypothetical protein MGC3079
 hypothetical protein FLJ10074
 protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)
 p30 DBC protein
 zinc finger protein 144 (Mei-18)
 BTG family, member 2
 cytochrome P450, subfamily XIB (steroid 11-beta-hydroxylase), polypeptide 1
 bone morphogenetic protein 1
 urotensin 2
 proteoglycan 4, (megakaryocyte stimulating factor, articular superficial zone protein, camptodactyly, arthropathy, coxa vara, pericarditis syndrome)
 quiescin Q6
 protein lysine phosphatase-like (proline instead of catalytic arginine), member a
 olfactory receptor, family 2, subfamily B, member 6
 Consensus includes gb:BE46525 /FEA=EST /DB_XREF=gi:9512223 /DB_XREF=esth94b10.x1 /CLONE=IMAGE:3195451 /UG=Hs.234773 Homo sapiens cDNA: FLJ22281
 KIAA1859 protein
 gamma-glutamyltransferase 1
 mitochondrial ribosomal protein S33
 desmoglein 3 (pemphigus vulgaris antigen)
 hypothetical protein FLJ22471
 huntingtin interacting protein 1
 acylphosphatase 2, muscle type
 KIAA0899 protein
 hypothetical protein FLJ20539
 hypothetical protein PRO0899
 VAMP (vesicle-associated membrane protein)-associated protein B and C
 eis variant gene 5 (eis-related molecule)
 core-binding factor, beta subunit
 hypothetical protein FLJ20399
 cyclin-dependent kinase 6
 Agt3, S. cerevisiae, homolog of
 phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila)
 carbonic anhydrase VB, mitochondrial
 retinoid X receptor, beta
 testis zinc finger protein
 solute carrier family 21 (organic anion transporter), member 9
 Consensus includes gb:AK025097.1 /DEF=Homo sapiens cDNA: FLJ21444 fls, clone COL04437, /FEA=mRNA /DB_XREF=gi:10437544 /UG=Hs.306764 Homo sapiens cDNA: AK025097
 aquaporin 6, kidney specific
 FAST thase
 gp25L2 protein
 cysteine and glycine-rich protein 1
 MCT-1 protein
 protein lysine phosphatase, non-receptor type 18 (brain-derived)
 receptor-interacting serine-threonine kinase 2
 hypothetical protein PRO1580
 solute carrier family 2 (facilitated glucose transporter), member 4
 protective protein for beta-galactosidase (galactosialdosis)
 hypothetical protein FLJ21799
 KIAA0435 gene product
 retinoic acid repressible protein
 homolog of mouse quaking OKI (KH domain RNA binding protein)
 degenerative spermatocyte homolog, lipid desaturase (Drosophila)
 protein phosphatase 4 regulatory subunit 2
 cytochrome b5 reductase 1 (BR9.1)
 Consensus includes gb:W74494 /FEA=EST /DB_XREF=gi:1384781 /DB_XREF=estizd75f10.s1 /CLONE=IMAGE:346507 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete W74494
 Consensus includes gb:AW500220 /FEA=EST /DB_XREF=gi:7112628 /DB_XREF=estUJ-HF-BND-akd-c02-0-U1.r1 /CLONE=IMAGE:3076610 /UG=Hs.332496 Human DNA ss AW500220

77

Consensus includes gb:BF877404 /FEA=EST /DB_XREF=gi:11951299 /DB_XREF=est:602087150F1 /CLONE=IMAGE-4251205 /UG=Hs.296356 Homo sapiens mRNA, cDNA	BF877404	1,14
kinase insert domain receptor (a type III receptor tyrosine kinase)	NM_002253	1,14
phosphatidylinositol-4-phosphate 5-kinase, type I, beta	U76561	1,14
CD83 antigen (activated B lymphocytes, immunoglobulin superfamily)	NM_004233	1,14
TC037 homolog	AF123539	1,14
cytochrome c oxidase subunit VIIa polypeptide 2 (liver)	NM_001885	1,14
hypothetical protein FLJ10402	NM_018085	1,14
dydrobrexins, alpha	U46746	1,14
cytosolic ovarian carcinoma antigen 1	S72904	1,14
estrogen receptor 2 (ER beta)	A8006590	1,14
E2F transcription factor 4, p107/p130-binding	S75174	1,14
Consensus includes gb:U79276.1 /DEF=Human clone 23960 mRNA sequence, /FEA=mRNA /DB_XREF=gi:1710244 /UG=Hs.307238 Human clone 23960 mRNA sequence	U79276	1,14
homologous to yeast nitrogen permease (candidate tumor suppressor)	NM_006545	1,14
small nuclear ribonucleoprotein polypeptide A'	NM_003090	1,14
pyruvate dehydrogenase kinase, isoenzyme 1	NM_002610	1,14
SOM DNA binding protein	XG3071	1,14
immunoglobulin lambda locus	AF047245	1,14
myeloid differentiation primary response gene (88)	U70451	1,14
hyposphosphatase I	AF077198	1,14
tissue specific transplantation antigen P35B	U58766	1,14
POU domain, class 3, transcription factor 2	NM_005604	1,14
H2B histone family, member J	NM_003524	1,14
amyloid beta precursor protein binding protein 1, S9K0	NM_003905	1,14
hypothetical protein FLJ13149	NM_021826	1,14
spectrin, beta, non-erythrocytic 1	N92501	1,14
ataxia telangiectasia mutated (includes complementation groups A, C and D)	U82828	1,14
G protein-coupled receptor 51	AF089755	1,14
myosin, heavy polypeptide 8, cardiac muscle, alpha (cardiomyopathy, hypertrophic 1)	D00943	1,14
tumor rejection antigen (gp96) 1	NM_003289	1,14
zinc finger protein 137 (clone pH2-30)	NM_003438	1,14
SEC23 related gene family, member A (S. cerevisiae)	BE643231	1,14
VAMP (vesicle-associated membrane protein)-associated protein A (33KD)	AK000918	1,14
hypothetical protein FLJ20604	NM_017897	1,14
WD40 protein C1a01	NM_004804	1,14
KIAA0395 protein	AK022844	1,14
TEA domain family member 4	NM_015082	1,14
hypothetical protein MGCS347	U63524	1,14
IQ motif containing GTPase activating protein 1	NM_024063	1,14
calcyl-phosphate memosyltransferase polypeptide 1, catalytic subunit	A1678073	1,14
HOCMA18P protein	NM_003859	1,14
HIV-1 rev binding protein 2	AA160161	1,14
triggering receptor expressed on myeloid cells 1	NM_007043	1,14
apoptotic protease activating factor	NM_018643	1,14
Consensus includes gb:AL122122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L098 (from clone DKFZp434L098), /FEA=mRNA /DB_XREF=gi:6102949 /UG=Hs.274559 H	AL122122	1,14
inositol polyphosphate-3-phosphatase, 40KD	AF149794	1,14
hypothetical protein	NM_005539	1,14
BRCA1 associated protein	AA046752	1,14
LIM domain-containing preferred translocation partner in lipoma	AF035620	1,14
microtubule-associated protein tau	BF221852	1,14
small nuclear ribonucleoprotein polypeptide B"	A1870749	1,14
mitochondrial ribosomal protein S14	NM_003092	1,14
gb:AF132204.1 /DEF=Homo sapiens PRO2259 mRNA, complete cds, /FEA=mRNA /PROD=PRO2259 /DB_XREF=gi:11493553 /UG=Hs.302159 Homo sapiens PRO2259 mRNA	BC254653	1,14
DKFZP727M1111 protein	NM_015540	1,14
hypothetical protein FLJ13648	NM_024584	1,14
regulatory solute carrier protein, family 1, member 1	AJ268381	1,14
KIAA0984 protein	BF346592	1,14
Consensus includes gb:AK000864.1 /DEF=Homo sapiens cDNA FLJ10002 fts, clone HEMBA1000046, /FEA=mRNA /DB_XREF=gi:7021188 /UG=Hs.296522 Homo sapiens cDNA AK000864	AK000864	1,14
alpha-2-glycoprotein 1, zinc	AC004522	1,14
GTF2i repeat domain-containing 1	NM_016328	1,14
hypothetical protein FLJ23514	NM_021627	1,14
CGI-43 protein	AL550875	1,14
glioma tumor suppressor candidate region gene 2	NM_015710	1,14

like mouse brain protein E48
Consensus includes gb:BE25050 /FEA=EST /DB_XREF=gi:129548 /DB_XREF=est:60110055F1 /CLONE=IMAGE:3350965 /UG=Hs.307137 Human DNA sequence from cl BE25050
inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta
hypothetical protein FLJ20069
ELK1, member of ETS oncogene family
gb:NM_025156.1 /DEF=Homo sapiens hypothetical protein FLJ12474 (FLJ12474), mRNA, /FEA=mRNA /GEN=FLJ12474 /PROD=hypothetical protein FLJ12474 /DB_XREF=gi:NM_025156
T cell receptor beta locus
SP110 nuclear body protein
hypothetical protein PRO2389
polyoma 1
carboxypeptidase D
hypothetical protein FLJ22672
KIA0150 protein
GSS953 protein
DEADM (Asp-Glu-Ala-AspHis) box binding protein 1
Integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDP2, MSK12)
synovial sarcoma, X breakpoint 3
bunyophilin, subfamily 2, member A2
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 7
similar to rat myomegalin
hypothetical protein FLJ13556
cleavage stimulation factor, 3' pre-RNA, subunit 2, 64KD
niflurin 2
Consensus includes gb:AW269792 /FEA=EST /DB_XREF=gi:6656822 /DB_XREF=est:tx45d03.x1 /CLONE=IMAGE:2816069 /UG=Hs.5010 Homo sapiens clone 24672 mRNA AW269792
poly(A)-specific ribonuclease (deadenylation nuclease)
KIA00556 protein
Consensus includes gb:BG491393 /FEA=EST /DB_XREF=gi:13452905 /DB_XREF=est:602535682F1 /CLONE=IMAGE:4684748 /UG=Hs.183110 EST's, Weakly similar to ALU BG491393
Integrin beta 3 binding protein (beta3-endonexin)
TNF receptor-associated factor 3
hypothetical gene MGC19595
Consensus includes gb:AK022065.1 /DEF=Homo sapiens cDNA FLJ12003 fs, clone HEMBB1001537 /FEA=mRNA /DB_XREF=gi:10433385 /UG=Hs.295683 Homo sapiens c AK022065
centaurin, gamma 1
HSPC016 protein
hypothetical protein FLJ13134
DiGeorge syndrome gene D
ubiquitin-conjugating enzyme E2D 2 (UBCA5 homolog, yeast)
RacQ protein-like 5
KIA0728 protein
low density lipoprotein receptor-related protein 5
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2
SAM domain and HD domain, 1
Consensus includes gb:AK024543.1 /DEF=Homo sapiens cDNA: FLJ20890 fs, clone ADKA03323 /FEA=mRNA /DB_XREF=gi:10436848 /UG=Hs.306586 Homo sapiens cDNA: AK024543
soluble carrier family 25 (mitochondrial carrier, brain), member 14
mitogen-activated protein kinase kinase kinase 5
gb:L14456.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-region, complete cds, /FEA=mRNA /DB_XREF=gi:398497 /FL=gb:L14456.1
hypothetical protein FLJ11787
ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
endothelin 2
polymerase (DNA directed), alpha
mitogen-activated protein kinase-activated protein kinase 2
plectin 1, intermediate filament binding protein, 500KD
retinoblastoma binding protein 6
KIAA1041 protein
hemoglobin, epsilon 1
transmembrane protein 4
hypothetical protein MGC2470
Consensus includes gb:AU147166 /FEA=EST /DB_XREF=gi:1008687 /DB_XREF=est:AU147166 /CLONE=HEMWB1002582 /UG=Hs.301683 Homo sapiens cDNA FLJ12091
BRF1 homolog, subunit of RNA polymerase II (transcription initiation factor IIB (S.cerevisiae))
guanine nucleotide binding protein (G protein), q polypeptide
transketolase-like 1
Slush-interacting protein
gamma-aminobutyric acid (GABA) A receptor, alpha 6
misato
NM_018116

MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	AI859865	1,14
sn3-associated polypeptide, 30kD	NM_003864	1,14
astroctadin 2	AK024064	1,14
potassium voltage-gated channel, Shaw-related subfamily, member 1	NM_004876	1,14
hypothetical protein MGC5149	U79260	1,14
KIAA0882 protein	BE646554	1,14
polymerase (RNA) II (DNA directed) polypeptide L (7.6kD)	AA772747	1,14
hypothetical protein FLJ10874	NM_018252	1,14
tailed-like kinase 2	AU151689	1,14
KIAA0494 gene product	NM_014774	1,14
hypothetical protein	BC001830	1,14
KIAA0121 gene product	D50911	1,14
myosin, light polypeptide 4, alkali; atrial, embryonic	AI133353	1,14
hepatocyte nuclear factor 4, alpha	NM_000457	1,14
DKFZP96G011 protein	AK023204	1,14
complement component 4 binding protein, alpha	NM_000715	1,14
hypothetical protein	BC001393	1,14
RNA binding protein S1, serine-rich domain	BC001659	1,14
reticulon 2	NM_005619	1,14
DnaJ (Hsp40) homolog, subfamily C, member 3	NM_006260	1,14
hypothetical protein	NM_016625	1,14
ribosomal protein L28 pseudogene 1	AW003989	1,14
UDP-glucose ceramide glucosyltransferase	NM_018093	1,14
GS2 gene	NM_003358	1,14
tripartite motif-containing 3	AI814551	1,13
NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1 (8kD, SDAP)	AA19307	1,13
butyrophilin, subfamily 2, member A2	NM_005003	1,13
NADPH oxidase 3	W58757	1,13
HIV-1 Rev binding protein	NM_015718	1,13
phospholamban	AI742626	1,13
Consensus includes gb:AK027161.1 /DEF=Homo sapiens cDNA: FLJ23508 fs, clone LNG03158, /FEA=mRNA /DB_XREF=gi:10440220 /UG=Hs.288772 Homo sapiens cDNA: AK027161	NM_002687	1,13
synaptotagmin 2	AF138616	1,13
phosphatidylinositol glycan, class C, pseudogene 1	AB000359	1,13
hypothetical protein FLJ14827	BE966197	1,13
hypothetical protein similar to beta-transducin family	NM_018096	1,13
supervillin	AW872377	1,13
polymerase (RNA) mitochondrial (DNA directed)	NM_005035	1,13
eukaryotic translation initiation factor 4 gamma, 3	AI768122	1,13
protease, serine, 2 (trypsin 2)	NM_002770	1,13
hemoglobin, gamma A	NM_000559	1,13
cerebroside (3'-phosphoadenylylsulfate:galactosylceramide 3') sulfotransferase	NM_004861	1,13
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8)	BC003874	1,13
glycophorin C (Gerbach blood group)	NM_002101	1,13
specie-type POZ protein	NM_003563	1,13
heterogeneous nuclear ribonucleoprotein R	BC001449	1,13
KIAA0414 protein	AI745225	1,13
hypothetical protein FLJ13441	NM_023924	1,13
tetratricopeptide repeat domain 3	D83077	1,13
nicastin	AF240468	1,13
galenin receptor 2	AF080586	1,13
lymphocyte antigen 117	AV713720	1,13
hypothetical protein FLJ20011	BC003086	1,13
hypothetical protein FLJ10477	NM_018105	1,13
G protein-coupled receptor kinase-interactor 2	AF124491	1,13
Consensus includes gb:AI653960 /FEA=EST /DB_XREF=gi:4737939 /DB_XREF=est:ty03h10.x1 /CLONE=IMAGE:2278051 /UG=Hs.196555 ESTs	AI653960	1,13
putative glycolipid transfer protein	NM_015899	1,13
TRK-like gene	NM_008070	1,13
Rhesus blood group, D antigen	L08429	1,13
capping protein (actin filament) muscle Z-line, beta	AL572341	1,13
ARP1 actin-related protein 1 homolog A, centractin alpha (yeast)	AL532341	1,13
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1	AA593983	1,13
zinc finger protein 144 (Mei-18)	AI744229	1,13

KIAA1030 protein	AB028953	1.13
platelet-derived growth factor receptor-like	NM_008207	1.13
glioma tumor suppressor candidate region gene 1	NM_015711	1.13
potassium voltage-gated channel, subfamily G, member 2	NM_012283	1.13
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6	NM_001685	1.13
growth associated protein 43	AF279774	1.13
melan-A	NM_005511	1.13
apoptosis regulator	NM_018561	1.13
hypothetical protein FLJ13490	NM_024942	1.13
heterogeneous nuclear ribonucleoprotein H3 (2H9)	AF132363	1.13
nuclear receptor coactivator 3	AA38999	1.13
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 3	NM_003653	1.13
Cluster Ind. A1832248:14910.1 Homo sapiens cDNA, 3' end /clone=IMAGE-2075682 /clone_end=3' /gp=A1832249 /ug=5452920 /ug=Hs.211522 /len=545	A1832249	1.13
KIAA0544 protein	AB011116	1.13
Cytidine-rich, angiogenic inducer, 61	AF003114	1.13
CD59 antigen p16-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EJ32 and G344)	AL513759	1.13
hypothetical protein FLJ22195	AL523965	1.13
cystatin 8 (cystatin-related epidermal specific)	NM_005492	1.13
hypothetical protein DKFZ664O043	AK001389	1.13
dUTP pyrophosphatase	AB049113	1.13
hypothetical protein FLJ14124	NM_024868	1.13
hemopexin	BC005395	1.13
Cyclin G associated kinase	D88435	1.13
down-regulator of transcription 1, TBP-binding (negative cofactor 2)	NM_001938	1.13
IRB (bacterial acetoacetate synthase)-like	AC004794	1.13
RAB27A, member RAS oncogene family	AF125393	1.13
Interleukin 19	NM_013371	1.13
spolipoprotein A-I	X02162	1.13
KIAA1277 protein	AA127623	1.13
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)	X55005	1.13
cystine-rich protein 2	U36190	1.13
A kinase (PRKA) anchor protein 1	U34074	1.13
PTD18 protein	NM_016125	1.13
ADP-ribosylation factor 3	NM_001659	1.13
retinoblastoma binding protein 7	NM_002893	1.13
KIAA1282 protein	AB033088	1.13
v-erb sarcoma virus CT10 oncogene homolog (avian)	NM_016823	1.13
KIAA0953 protein	AB023170	1.13
Tax interaction protein 1	AK001327	1.13
amphiphysin (Slit-Mann syndrome with breast cancer 128kD autoantigen)	NM_001635	1.13
associated molecule with the SH3 domain of STAM	NM_006463	1.13
GLUT4 enhancer factor	NM_020062	1.13
v-ets erythroblastosis virus E26 oncogene homolog 1 (avian)	NM_005238	1.13
solute carrier family 21 (organic anion transporter), member 3	AF085224	1.13
hypothetical protein FLJ23233	AA429615	1.13
KIAA0853 protein	AL136745	1.13
sorting nexin 3	BE962615	1.13
T-box 21	NM_013351	1.13
heterogeneous nuclear ribonucleoprotein O (AU-rich element RNA binding protein 1, 37kD)	W74620	1.13
phosphoribosyl pyrophosphate synthetase-associated protein 1	NM_002766	1.13
growth hormone inducible transmembrane protein	AL136713	1.13
keratin, hair, acidic38	NM_002279	1.13
hypothetical protein FLJ11106	NM_018324	1.13
potassium voltage-gated channel, KQT-like subfamily, member 2	NM_004516	1.13
calcium/calmodulin-dependent protein kinase (CaM kinase) II beta	U23460	1.13
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	NM_003070	1.13
Consensus includes gb-B03390664 /FEA=EST /DB_XREF=gi:13284112 /DB_XREF=est:602415258 / /CLONE=IMAGE-4523487 /UG=Hs.97496 YY1 transcription factor /FL=gb-B03390664	BC3390664	1.13
neutrophin 3	NM_002527	1.13
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2	AL542253	1.13
Rho-associated, coiled-coil containing protein kinase 1	N22548	1.13
cystatin S	NM_001859	1.13
mammalian-binding lectin serine protease 2	AB008047	1.13
BCL2-antagonist of cell death	U66879	1.13

KIAA0268 protein
 GA binding protein transcription factor, alpha subunit (80kD)
 calcium channel, voltage-dependent, alpha 2/delta subunit 1
 zinc finger protein 302
 neuroblastoma, suppression of tumorigenicity 1
 general transcription factor IIC, polypeptide 1 (alpha subunit, 220kD)
 serine/threonine kinase 17a (apoptosis-inducing)
 serine palmitoyltransferase, long chain base subunit 2
 HIV-1 inducer of short transcript binding protein; lymphoma related factor
 HEMK homolog 7/b
 arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)
 KIAA0494 gene product
 hypothetical gene supported by AF038182; BC009203
 growth arrest-specific 6
 protection of telomeres 1
 transposon-derived Butler1 transposase-like protein
 FH1/FH2 domain-containing protein
 RAN binding protein 17
 CGI-74 protein
 BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like
 hypothetical protein MGC22654
 Consensus includes gb:AL050002.1 /DEF=Homo sapiens mRNA, cDNA DKFZp564O222 (from clone DKFZp564O222). /FEA=mRNA /DB_XREF=gi:4884256 /UG=Hs.94795 H-
 ARP2 actin-related protein 2 homolog (yeast)
 putative membrane protein
 decalin
 CCH3 general control of amino-acid synthesis 5-like 2 (yeast)
 Consensus includes gb:A1923972 /FEA=EST /DB_XREF=gi:5659936 /DB_XREF=est:wn49c10.x1 /CLONE=IMAGE2448786 /UG=Hs.127723 Homo sapiens mRNA; cDNA DKFZp564O222
 hypothetical protein R30953_1
 chromosome 17 open reading frame 1A
 Wiskott-Aldrich syndrome protein interacting protein
 Consensus includes gb:AK000270.1 /DEF=Homo sapiens cDNA FLJ20263 fs, clone COLF7804, highly similar to AJ131693 Homo sapiens mRNA for AXAP450 protein. /FEA=AK000270
 interferon, gamma
 KIAA1069 protein
 mesothionein 1X
 KIAA0915 gene product
 hypothetical protein FLJ10468
 platophilin 4
 ubiquitin-activating enzyme E1-like
 hypothetical protein
 KIAA0409 protein
 Mees1, myeloid ecotropic viral integration site 1 homolog 2 (mouse)
 calpain 7
 RAD21 homolog (S. pombe)
 hypothetical protein
 CCAA1/enhancer binding protein (CESBP), delta
 Consensus includes gb:Z98950 /DEF=Human DNA sequence from PAC 507115 on chromosome Xq28.3-27.3. Contains 605 ribosomal protein L44 (L41, L36) like gene, ESTs, Z98950
 PTPRF interacting protein, binding protein 1 (liprin beta 1)
 cadherin 16, XSP-cadherin
 Kruppel-like factor 4 (gut)
 calcium channel, voltage-dependent, alpha 1 subunit
 hypothetical protein FLJ12496
 integrin, alpha 4 (enigen CD49D, alpha 4 subunit of VLA-4 receptor)
 mesh homeo box homolog 2 (Drosophila)
 laminin, beta 3 (lncin (125kD), lamin (140kD), BM600 (125kD))
 phosphatidylinositol-3-phosphate 3-phosphatase adaptor subunit
 hypothetical protein FLJ22222
 casein kinase 1, alpha 1
 interferon 1, alpha
 CD58 antigen, (lymphocyte function-associated antigen 3)
 HNOEL-iso protein
 U8 snRNA-associated Sm-like protein
 synapobin 2
 Consensus includes gb:AK022679.1 /DEF=Homo sapiens cDNA FLJ12617 fs, clone NT2RM4001662. /FEA=mRNA /DB_XREF=gi:10434213 /UG=Hs.287538 Homo sapiens c AK022679

CD28 antigen (Tp44)
 ferritin, heavy polypeptide 1
 myosin 5C
 HUG1 gene
 ribosomal protein S29
 glycosylphosphatidylinositol specific phospholipase D1
 Consensus includes gp:AL049037 /FEA=EST /DB_XREF=gi:4728346 /DB_XREF=est:DKFZp343P1518_r1 /CLONE=DKFZp343P1518 /UG=Hs.323833 syntaphilin
 PR domain containing 4
 hypothetical C2H2 zinc finger protein FLJ22504
 hypothetical protein FLJ20150
 Consensus includes gp:AK021982.1 /DEF=Homo sapiens cDNA FLJ11930 fls, clone HEMBB 100044.1. /FEA=mRNA /DB_XREF=gi:0433501 /UG=Hs.306618 Homo sapiens c AK021992
 beaded filament structural protein 2, p14kin
 hypothetical protein MQC2821
 Consensus includes gp:AL049037 /FEA=EST /DB_XREF=gi:5425467 /DB_XREF=est:w70g04.x1 /CLONE=IMAGE:2408214 /UG=Hs.194051 Homo sapiens mRNA; cDNA DKF A1814252
 leukotriene b4 receptor (chemokine receptor-like 1)
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 7
 perlecanin material 1
 P311 protein
 hypothetical protein FLJ12438
 RAN binding protein 2
 synovial sarcoma translocation, chromosome 18
 15 kDa selenoprotein
 tyrosine kinase, non-receptor, 1
 core promoter element binding protein
 hypothetical protein FLJ11132
 RNA binding motif protein 15
 chondroitin sulfate proteoglycan 4 (melanoma-associated)
 TIRAM-like protein
 hypothetical protein DKFZp343E2135
 CD2 antigen (cytoplasmic tail) binding protein 2
 hypothetical protein MQC2508
 protein tyrosine phosphatase type IVA, member 3
 retinoblastoma (X-linked, juvenile) 1
 methylmalonate-semialdehyde dehydrogenase
 mitogen-activated protein kinase 3
 apolipoprotein E
 TEA domain family member 1 (SV40 transcriptional enhancer factor)
 farnesyltransferase, CAAX box, alpha
 ubiquitin-conjugating enzyme E2C
 spermatogenesis associated 2
 hypothetical protein MQC4126
 solute carrier family 19 (ololate transporter), member 1
 ribosomal protein S20
 Consensus includes gb:D16471.1 /DEF=Human mRNA, Xq terminal portion. /FEA=mRNA /DB_XREF=gi:495089 /UG=Hs.121571 Human mRNA, Xq terminal portion
 vascular endothelial growth factor
 pyridine-5-carboxylate reductase 1
 homeo box 11-like 2
 hypothetical protein MQC4178
 cyclinomatosis (turban tumor syndrome)
 40S ribosomal protein S27 isoform
 growth differentiation factor 11
 wingless-type MMTV integrallon site family, member 6
 opoid receptor, kappa 1
 endocytic receptor (macrophage mannose receptor family)
 SHG-domain binding protein 2
 glioma amplified on chromosome 1 protein (leucine-rich)
 KIAA0733 protein
 cyclic nucleotide gated channel beta 3
 vitellin-associated protein VIT-1
 phospholipase C, epsilon 2
 transmembrane 4 superfamily member 8
 Consensus includes gb:AL023584 /DEF=Human DNA sequence from clone 67K17 on chromosome 6q24.1-24.3. Contains the HIVEP2 (Schnurri-2) gene for HIV type 1 Enhanc
 M10098 Human 18S rRNA sequence, length 1869 bases, 5 prime target bases 1-646

AF222341 1.12
 AA083483 1.12
 NM_018728 1.12
 NM_020527 1.12
 NM_001032 1.12
 AL031230 1.12
 AL049037 1.12
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 NM_022095 1.12
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 NM_000912 1.12
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 NM_019088 1.11
 AL117620 1.11
 AL117515 1.11
 AF053453 1.11
 AF023584 1.11
 M10098 1.11

84

hypothetical protein DKFZ0434.037 NM_030952 1.11
Consensus includes gb:AA148507 /FEA=EST /DB_XREF=gi:1721551 /DB_XREF=est:z06005.s1 /CLONE=IMAGE:491528 /UG=Hs.14846 Homo sapiens mRNA; cDNA DKFZ: AA148507 1.11
zinc finger protein 157 (HZF22) NM_003446 1.11
lactin (Strongylocentrotus purpuratus) homolog 2 (actin-bundling protein, retinal) NM_012418 1.11
golgi phosphoprotein 5 AK001574 1.11
tyrosine phosphatase II NM_007260 1.11
zinc finger protein NM_014347 1.11
progesterone-induced blocking factor 1 NM_006346 1.11
kinesin-like 3 NM_005355 1.11
POU domain, class 5, transcription factor 1 AF268617 1.11
homo box D1 A1168371 1.11
Consensus includes gb:AU143746 /FEA=EST /DB_XREF=gi:11007267 /DB_XREF=est:AU143746 /CLONE=HEMBA1005579 /UG=Hs.82193 esterase Diormylglutathione hydrolase 1.11
ATPase, vacuolar, 14 kD NM_004231 1.11
rabi3 GTPase-activating protein, non-catalytic subunit (150kD) NM_012414 1.11
growth hormone 1 BF223718 1.11
tubule functional domain (PTPRF Interacting) NM_000515 1.11
inner membrane protein, mitochondrial (mitofilin) NM_006839 1.11
solute carrier family 2 (facilitated glucose/fructose transporter), member 5 NM_003039 1.11
cysteine binding protein 2 NM_030758 1.11
Consensus includes gb:AL078598 DEF=Human DNA sequence from clone RP3-42905 on chromosome 6q21-22.1. Contains the NR2E1 gene for nuclear receptor 2E1 (rat) 1.11
phosphodiesterase 2A, cGMP-stimulated NM_002599 1.11
captein 3, (p94) AF127765 1.11
allograft inflammatory factor 1 BF213829 1.11
interferon regulatory factor 4 D78261 1.11
GA binding protein transcription factor, beta subunit 2 (47kD) NM_002041 1.11
for protein disulfide isomerase-related NM_006810 1.11
tumor protein 63 kDa with strong homology to p53 AB010153 1.11
hypothetical protein FLJ12270 NM_024673 1.11
tumor necrosis factor receptor superfamily, member 1A NM_001065 1.11
immunoglobulin heavy constant gamma 3 (G3m marker) M87789 1.11
guanine nucleotide binding protein (G protein), gamma 7 BF432795 1.11
tumor necrosis factor, alpha-induced protein 3 A1736896 1.11
Consensus includes gb:AK024135.1 /DEF=Homo sapiens cDNA FLJ14073 fts, clone HEMBB1001812 /FEA=mRNA /DB_XREF=gi:10436441 /UG=Hs.296751 Homo sapiens 1.11
v-ort avian sarcoma virus CT10 oncogene homolog NM_003931 1.11
WAS protein family, member 1 NM_019853 1.11
protein phosphatase 4 regulatory subunit 2 A4339556 1.11
thionotin interacting protein A9235734 1.11
Consensus includes gb:AB25734 /FEA=EST /DB_XREF=gi:5661698 /DB_XREF=est:wo34q08.x1 /CLONE=IMAGE:2457278 /UG=Hs.110613 KIAA00220 protein 1.11
polymerase (RNA) III (DNA directed) (39kD) NM_006466 1.11
calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma A0933569 1.11
KIAA1018 protein NM_014967 1.11
CD22 antigen NM_024064 1.11
F-box only protein 2 NM_001771 1.11
homo box C11 NM_012168 1.11
HSPC255 protein NM_014212 1.11
hypothetical protein PP1044 NM_014153 1.11
baculoviral IAP repeat-containing 3 NM_021730 1.11
neutral sphingomyelinase (N-SMase) activation associated factor U37546 1.11
phosphodiesterase 4D interacting protein (myomegalin) NM_003580 1.11
small inducible cytokine A1, I-309 AB042555 1.11
triple helix repeat containing 4 NM_002981 1.11
crystallin, zeta (quinone reductase)-like 1 NM_007185 1.11
hypothetical protein FLJ10700 NM_005111 1.11
KIAA0009 gene product NM_018182 1.11
leiomodin 1 (smooth muscle) BF214329 1.11
activating transcription factor 6 BC001755 1.11
mitochondrial ribosomal protein L13 NM_007348 1.11
ubiquitin-conjugating enzyme E2A (RAD6 homolog) NM_014078 1.11
interferon, gamma-inducible protein 16 NM_003336 1.11
mitochondrial ribosomal protein S22 BG256877 1.11
CGI-130 protein NM_020191 1.11
Consensus includes gb:AW274747 /FEA=EST /DB_XREF=gi:5661777 /DB_XREF=est:sn34e08.x1 /CLONE=IMAGE:2695622 /UG=Hs.270527 Homo sapiens mRNA; cDNA Df: AW274747 1.11

hypothetical protein ASH1
Consensus includes gb:AU144284 /FEA=EST /DB_XREF=gi:1005805 /DB_XREF=est:AU144284 /CLONE=HEMBA1001439 /UG=Hs.11801 interferon regulatory factor 6 /FL= AU144284
transformation/transcription domain-associated protein
brain link protein-1
Consensus includes gb:AK021980.1 /DEF=Homo sapiens cDNA FLJ11898 fs. clone HEMBA1007322. /FEA=mRNA /DB_XREF=gi:10433269 /UG=Hs.313509 Homo sapiens c
translocase of outer mitochondrial membrane 70 homolog A (yeast)
KIAA1089 protein
CDC10 cell division cycle 10 homolog (S. cerevisiae)
hypothetical protein
Consensus includes gb:BE549732 /FEA=EST /DB_XREF=gi:9791424 /DB_XREF=est:7638a04.x1 /CLONE=IMAGE:3230478 /UG=Hs.112158 Homo sapiens PAC clone RP4-7 BE549732
upholipase kinase type 2 isoform
gb:BC002361.1 /DEF=Homo sapiens. Similar to transducin (beta)-like 3. clone MGC:8613. mRNA, complete cds. /FEA=mRNA /PROD=Similar to transducin (beta)-like 3 /DB_XREF=gi:10433269 /UG=Hs.313509 Homo sapiens PAC clone RP4-7 BE549732
SHP2 interacting transmembrane adaptor
centaurin, delta 1
KIAA1033 protein
serine palmitoyltransferase, long chain base subunit 1
gelosin (amyloidosis, Finnish type)
hypothetical protein FLJ10233
major histocompatibility complex, class II, DP beta 1
DKFZPS47E1010 protein
RNA binding motif, single stranded interacting protein 1
DKFZP434H132 protein
Consensus includes gb:AW873544 /FEA=EST /DB_XREF=gi:8007597 /DB_XREF=est:h62a01.x1 /CLONE=IMAGE:3041928 /UG=Hs.277500 Human mRNA for T cell recepto
hypothetical protein FLJ11099
metallothionein 1L
prostaglandin E receptor 4 (subtype EP4)
aridine homolog 2 (Drosophila)
transcription factor-like 4
ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) 9kD
chromosome 18 open reading frame 1
leucine rich protein
nucleophosmin (nuclear phosphoprotein B23, numatrin)
KIN, antigenic determinant of recA protein homolog (mouse)
palate, lung and nasal epithelium carcinoma associated
phospholipase A2, group IVB (cytosolic)
KIAA0170 gene product
transcription elongation factor B (SIII), polypeptide 1-like
trypsin-like peptidase II
Sac domain-containing inositol phosphatase 2
Consensus includes gb:AK000787.1 /DEF=Homo sapiens cDNA FLJ20780 fs. clone COL04256. /FEA=mRNA /DB_XREF=gi:7021089 /UG=Hs.157392 Homo sapiens cDNA FI
fodrin 2
alpha-1-microglobulin/bikunin precursor
immunoglobulin kappa variable 1-13
gene from NF2/meringoma region of 22q12
methylmalonate-semialdehyde dehydrogenase
interleukin 5 receptor, alpha
mitogen-activated protein kinase kinase 1
forhead box O3A
baculoviral IAP repeat-containing 4
NADH dehydrogenase (ubiquinone) Fc-S protein 4 (18kD) (NADH-coenzyme Q reductase)
hypothetical protein FLJ10415
Consensus includes gb:AL080132.1 /DEF=Homo sapiens mRNA; cDNA DKFZP434F083 (from clone DKFZP434F083). /FEA=mRNA /DB_XREF=gi:5262572 /UG=Hs.118207 H
UDP-galactose transporter related
hypothetical protein DKFZP434I1916
trypsin, alpha
zinc finger protein 184 (Kruppel-like)
cytochrome b-581
heme oxygenase (decycling) 2
mitogen-activated protein kinase 4
LAG1 longevity assurance homolog 2 (S. cerevisiae)
KIAA0708 protein
c-myc binding protein
huntingtin (Huntington disease)

NM_018469
NM_003496
NM_021817
AK021980
BC003633
AB029012
AU157515
BC000979
AF245447
BC002361
NM_014450
AK023737
AK001728
AL568804
NM_000177
NM_018034
NM_002121
NM_015607
D28482
NM_015492
AW873544
NM_018320
NM_002450
AI675173
AW074630
BE984655
NM_003945
AI345006
NM_007244
BC002398
NM_012311
NM_016583
AK000550
NM_014641
BE984043
AW612376
NM_014937
AK000787
BC003683
NM_001633
AW408194
AB023200
NM_005589
NM_000584
AI571419
NM_001455
U32974
BC005270
NM_018089
H.AL080132
NM_005827
AL136771
NM_003293
AB111577
AL514271
D21243
BF115223
AK001105
AB014608
AL525412
NM_002111

somatostatin receptor 3	NM_001051	1.11
dictyopt homolog 4 (Xenopus laevis)	NM_014420	1.11
ephrin-B1	NM_004429	1.11
transcription factor 8 (represses interleukin 2 expression)	A086174	1.11
COQ7 coenzyme Q 7 homolog ubiquinone (yeast)	AF032900	1.11
Bacillus subtilis REF-L38424 /DEF-F8 subtilis dppB, jofF, jofG genes corresponding to nucleotides 2634-3089 of L38424 /LEN=1931 (-5, -M, -3) represent transcript regions 5 pL38424		1.11
Consensus includes gb:AF305057 /DEF=Homo sapiens RTS (RTS) gene, complete cds, alternatively spliced /FEA=mrRNA_1 /DB_XREF=gi:17094017 /UG=Hs.180433 rTS bet AF305057		1.11
ELG protein	NM_018553	1.11
CASP8 and FADD-like apoptosis regulator	AF041461	1.11
CD2-associated protein	NM_012120	1.11
branched chain keto acid dehydrogenase E1, beta polypeptide (maple syrup urine disease)	AL531533	1.11
SET domain, bifurcated 1	NM_012432	1.11
Consensus includes gb:AJ39732 /FEA=EST /DB_XREF=gi:4076659 /DB_XREF=est:qq42005.x1 /CLONE=IMAGE:1935201 /UG=Hs.309763 ESTs, Weakly similar to ALU7_HL AJ39732		1.10
chloride channel Ks	NM_004070	1.10
hypothetical protein FLJ23420	NM_025061	1.10
spastic paraplegia 4 (autosomal dominant; spasin)	A8078006	1.10
S100 calcium binding protein A1	NM_008271	1.10
transcription factor-like 5 (basic helix-loop-helix)	AL035689	1.10
DKFZP586G011 protein	AK021613	1.10
three prime repair exonuclease 1	BC002903	1.10
mitochondrial ribosomal protein S35	NM_021821	1.10
myelin basic protein	AW070431	1.10
tyrosyl-tRNA synthetase 2 (mitochondrial)	NM_015838	1.10
transmembrane 9 superfamily member 1	BE899402	1.10
ectonucleoside triphosphate diphosphohydrolase 1	AV717590	1.10
hydroxysteroid (17-beta) dehydrogenase 1	NM_000413	1.10
OnaJ (Hsp40) homolog, subfamily B, member 4	NM_007034	1.10
hypothetical protein FLJ12505	NM_024749	1.10
postrypsinic protein CRIPT	NM_014171	1.10
cyclochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 7, pseudogene 2	U22044	1.10
PCTAIRE protein kinase 1	BC001048	1.10
parminin 1	NM_015388	1.10
sema domain, immunoglobulin domain (Ig), transmembrane domain TM) and short cytoplasmic domain, (semaphorin) 4C	NM_017789	1.10
amyloid beta (A4) precursor-like protein 2	NM_001642	1.10
hypothetical protein DKFZp434O1427	NM_030963	1.10
HOG-like suppressor	NM_020386	1.10
mitogen-activated protein kinase-activated protein kinase 5	NM_003668	1.10
ring finger protein 24	NM_022823	1.10
transcription factor-like 5 (basic helix-loop-helix)	BC000213	1.10
KRAAB-zinc finger protein SZF 1-1	NM_008602	1.10
uracil-DNA glycosylase pseudogene 2	AF114817	1.10
v-yes-1 Yernaguchi sarcoma viral related oncogene homolog	X82986	1.10
frequently rearranged in advanced T-cell lymphomas		1.10
synaptin 4A (placental)	AI358412	1.10
lin D proto-oncogene	NM_005479	1.10
HCF-binding transcription factor Zhangfei	NM_004604	1.10
nucleoporm 62AD	NM_005354	1.10
Cluster Ind. AJ302185:qr58d02.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:1902435 /clone_end=3' /gb=AI302185 /gi=3961531 /ug=Hs.4789 /len=522	NM_012346	1.10
keratoprotein alpha 6 (keratin alpha 7)	AI002185	1.10
CD58 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	X16447	1.10
HSPC932 protein	NM_014150	1.10
KIAA0283 gene product	BE614199	1.10
microtubule-interacting protein that associates with TRAF3	AF230877	1.10
SEC23 protein	AK001465	1.10
dyspenia 1, longon (autosomal dominant; torsin A)	BC000674	1.10
expression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein)	U17714	1.10
tblod-like 1	AI769310	1.10
hypothetical protein FLJ10111	NM_017999	1.10
nuclear transcription factor Y, beta	AI804118	1.10
melanoma antigen, family B, 1	NM_002363	1.10
KIAA1668 protein	W46406	1.10
Consensus includes gb:AA133342 /FEA=EST /DB_XREF=gi:1690310 /DB_XREF=est:zn92h03.s1 /CLONE=IMAGE:585685 /UG=Hs.6700 ESTs	AA133342	1.10

hypothetical protein FLJ20450
 KIAA0931 protein
 HIRA interacting protein 3
 ferredoxin reductase
 hypothetical protein FLJ20644
 hypothetical protein FLJ20188
 cholinergic receptor, nicotinic, alpha polypeptide 2 (neuronal)
 KIAA1284 protein
 testis expressed sequence 13A
 nardilysin (N-angiotensin diester convertase)
 hypothetical protein FLJ10392
 gpA1072523.1 IDEF+Homo sapiens transcription factor T2P (T2P) mRNA, complete cds. /FEA=mRNA /GEN=T2P /PROD=transcription factor T2P /DB_XREF=gi:13195150 /A:AY027523
 set-1 suppressor of lin-12-like (C. elegans)
 collagen, type XVIII, alpha 1
 phosphatidylinositol transfer protein
 interleukin 16 (lymphocyte chemoattractant factor)
 proliferation-associated 204, 38kD
 Consensus includes gp:AL137398.1 IDEF+Homo sapiens mRNA, cDNA DKFZp434K052 (from clone DKFZp434K052). /FEA=mRNA /DB_XREF=gi:6807944 /UG+Hs.169639 /A:U137398
 lymphocyte antigen 17
 Consensus includes gp:AL137398.1 IDEF+Homo sapiens mRNA, cDNA DKFZp434K052 (from clone DKFZp434K052). /FEA=mRNA /DB_XREF=gi:6807944 /UG+Hs.169639 /A:U137398
 sialyltransferase 8 (alpha-2, 8-polysialyltransferase) D
 solute carrier family 6 (neurotransmitter transporter, L-proline), member 7
 hypothetical protein FLJ14827
 KIAA0767 protein
 nucleotide-sugar transporter similar to C. elegans sqv-7
 GTP-binding protein regB
 KIAA0916 protein
 jagged 1 (Alagille syndrome)
 chromosome 2 open reading frame 6
 dystrobrevin, alpha
 ref1004
 Consensus includes gp:A041204 /FEA=EST /DB_XREF=gi:3280398 /DB_XREF=est:ov77g06.x1 /CLONE=IMAGE:1643386 /UG+Hs.323748 Homo sapiens clone CDABP008c A041204
 solute carrier family 6 (neurotransmitter transporter, glycine), member 9
 breast carcinoma amplified sequence 4
 COI-02 protein
 ATP-binding cassette, sub-family C (CFTR/MRP), member 4
 hypothetical protein FLJ23445
 dishevelled, dish homolog 3 (Drosophila)
 xenoderm pigmentum, complementation group C
 regulator of G-protein signalling 1
 gp:NM_004692.1 IDEF+Homo sapiens internerxin neuronal intermediate filament protein, alpha (INA), mRNA. /FEA=mRNA /GEN=INA /PROD=intermedial neuronal intermediate filament protein, alpha (INA). /FEA=mRNA /DB_XREF=gi:897614 /FL=gb:L08961.1
 putative human HLA class II associated protein 1
 KIAA0578 protein
 suppressor of Ty 8 homolog (S. cerevisiae)
 solute carrier family 24 (sodium/potassium/calcium exchanger), member 1
 hypothetical protein DKFZp564A022
 cAMP responsive element modulator
 dimethylarginine dimethylaminohydrolase 2
 TNFRSF1A-associated via death domain
 LIM domain only 6
 gp:L08961.1 IDEF+Homo sapiens transmembrane tyrosine kinase mRNA, complete cds. /FEA=CDs /PROD=tyrosine kinase /DB_XREF=gi:897614 /FL=gb:L08961.1
 syntaxin 3A
 neogelin homolog 1 (chicken)
 BWRT protein
 X transporter protein 3
 mucosal vascular addressin cell adhesion molecule 1
 thyroid hormone receptor-associated protein, 95-kD subunit
 transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor
 hypothetical protein FLJ22055
 syntaxin binding protein 2
 baculoviral IAP repeat-containing 5 (sunvin)
 UBX domain-containing 1
 KIAA0483 protein

hypothetical protein FLJ14213	NM_024841	1.10
hypothetical protein FLJ23132	NM_025054	1.10
KIA0329 gene product	NM_014844	1.10
lifeguard	AB023167	1.10
hypothetical protein FLJ12671	AW284567	1.10
MYLE protein	NM_014015	1.10
tumor necrosis factor receptor superfamily, member 10b	AF016266	1.10
A kinase (PRKA) anchor protein 10	A4456929	1.10
nuclear respiratory factor 1	L22454	1.10
phosphoribosyl pyrophosphate synthetase 1	BC001605	1.10
Glutamate dehydrogenase-2	AC006144	1.10
SFRS protein kinase 2	AC005070	1.10
insulin receptor-related receptor	J05046	1.10
protein phosphatase	NM_016364	1.10
2'-5'-oligoadenylate synthetase 2 (69-71 kD)	NM_002535	1.10
NAADPH oxidase 1	NM_013955	1.10
iodine receptor 1	AL050262	1.10
dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS270, DXS272	NM_004010	1.10
actin, gamma 2, smooth muscle, enteric	NM_001615	1.10
Rag C protein	NM_022157	1.10
peptidylglycine alpha-amidating monooxygenase COOH-terminal interactor	AF056209	1.10
pre-mRNA splicing SR protein RA4	AA646521	1.10
Consensus includes gb:AF035278.1 /DEF+Homo sapiens clone 23676 mRNA sequence, /FEA=mRNA /DB_XREF=gi:2661027 /UG=Hs.100841 Homo sapiens clone 23676 mf	AF035278	1.10
SUMO-1 activating enzyme subunit 2	NM_004940	1.10
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 7 (RNA helicase, 52kD)	NM_004940	1.10
Consensus includes gb:AL049288.1 /DEF+Homo sapiens mRNA; cDNA DKFZp564G103 (from clone DKFZp564G103), /FEA=mRNA /DB_XREF=gi:4500017 /UG=Hs.2026841	AL049288	1.10
chromosome 11 open reading frame 13	NM_003475	1.10
sphingosine kinase type 2 isoform	AA483440	1.10
N-acetylglucosaminidase, alpha-	M38083	1.10
ADP-ribosylation factor-like 7	BC435404	1.10
Consensus includes gb:BE673445 /FEA+EST /DB_XREF=gi:10033966 /DB_XREF=est:7d35d10.x1 /CLONE=IMAGE:3249235 /UG=Hs.22049 Homo sapiens chromosome 19,	BE673445	1.10
protein kinase C-like 1	NM_007741	1.10
copine III	NM_003909	1.10
hypothetical protein FLJ12085	NM_022771	1.10
importin alpha 6 (importin alpha 7)	AU154321	1.10
nuclein domain 10 protein	BC004130	1.10
lectin 2 (testicular)	AB033823	1.10
p21(CDKN1A)-activated kinase 4	NM_005884	1.10
dystrophia myotonia-containing WD repeat motif	L19267	1.10
homo box B1	NM_002144	1.10
Consensus includes gb:A010895 /FEA+EST /DB_XREF=gi:3630631 /DB_XREF=est:wd20n09.x1 /CLONE=IMAGE:2328737 /UG=Hs.19720 Homo sapiens mRNA full length in	A010895	1.10
putative methyltransferase	AW069290	1.10
KIA02595 protein	AI674759	1.10
Consensus includes gb:A010195 /DEF+Homo sapiens ELK1 pseudogene (ELK2) and immunoglobulin heavy chain gamma pseudogene (IGHGP) /FEA=CDS /DB_XREF=gi:4	A010195	1.10
ATP-binding cassette, sub-family C (CFTRMRP), member 1	AI539710	1.10
dystrophin related protein 2	NM_001939	1.10
hypothetical protein MGC3731	AW002578	1.10
hypothetical protein FLJ21616	NM_024675	1.10
mitogen-activated protein kinase kinase kinase 1	NM_007161	1.10
SUMO-1-specific protease	AF306506	1.10
KIA00648 protein	AK021757	1.10
mitochondrial carrier family protein	NM_018843	1.10
hypothetical protein FLJ10647	BC006241	1.10
cardioembryonic antigen-related cell adhesion molecule 8	M33326	1.10
Down syndrome critical region gene 1-like 2	NM_013441	1.10
apoptosis-related protein PNAS-1	NM_016534	1.10
HUBA-inducible	NM_006460	1.10
dystrophia myotonia-protein kinase	L08835	1.09
Consensus includes gb:AL390738 /DEF+Human DNA sequence from clone RP11-438F9 on chromosome 13 Contains the gene for the ortholog of mouse homeobox protein G:	AL390738	1.09
chorionic gonadotropin hormone 2	NM_022645	1.09
low density lipoprotein receptor (familial hypercholesterolemia)	NM_000527	1.09
ubiquitin specific protease 9, X chromosome (Drosophila (fat facets related)	AA824366	1.09
Cas-Bi-M (murine) ecotropic retroviral transforming sequence	NM_005188	1.09

thrombosane A2 receptor
 major histocompatibility complex, class II, DM alpha
 NK2 transcription factor homolog B (Drosophila)
 Consensus includes gp:AW303476/FEA-EST/DB_XREF=est:xx19e10.x1/CLONE=IMAGE2813610/UG=Hs.132299 ESTs
 ghorylita reductase/hydroxypyruvate reductase
 testis expressed sequence 13B
 ATP-binding cassette, sub-family B (MDR/TAP), member 6
 collagen, type VI, alpha 1
 offactory receptor, family 2, subfamily N, member 1 pseudogene
 Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2
 hypothetical protein FLJ13119
 M11167 Human 28S rRNA sequence, length 5025 bases, middle target bases 1668-3330
 Consensus includes gbl:10374.1/DEF=Human (clone CTG-A4) mRNA sequence. /FEA=mRNA/DB_XREF=gj:307286/UG=Hs.281587 Human (clone CTG-A4) mRNA sequen
 G protein-coupled receptor, family C, group 5, member B
 endothelin type b receptor-like protein 2
 acute carrier family 4, sodium bicarbonate cotransporter, member 8
 TP53 target gene 1
 hypothetical protein FLJ10849
 cystatin F (leukocystatin)
 Consensus includes gb:A620911/FEA-EST/DB_XREF=gj:4630037/DB_XREF=est:u05412.x1/CLONE=IMAGE2250167/UG=Hs.205125 ESTs
 LIM protein (similar to rat protein kinase C-binding enzyme)
 gb:L21758.1/DEF=Homo sapiens acute myeloid leukemia associated protein (AML1EAP) mRNA, complete cds. /FEA=mRNA/GEN=AML1EAP/DB_XREF=gj:400340/FL=gb:L21758
 hypothetical protein FLJ20424
 KIAA0397 gene product
 growth hormone 1
 HMB-Inducible
 RAB3 GTPase-ACTIVATING PROTEIN
 allograft inflammatory factor 1
 peptidylprolyl isomerase E (cyclophilin E)
 MYC-associated zinc finger protein (purine-binding transcription factor)
 N-acetyltransferase 2 (erythrine N-acetyltransferase)
 hypothetical protein PRO0871
 hypothetical protein MGC11349
 T-cell, immune regulator 1
 sodium bicarbonate transporter 4
 a disintegrin and metalloproteinase domain 28
 rat finger protein 2
 Consensus includes gb:A8050195.1/DEF=Homo sapiens mRNA, partial cds, clone H2-5, MJD homolog. /FEA=mRNA/DB_XREF=gj:11559487/UG=Hs.302079 Homo sapiens
 phosphodiesterase 8A
 potassium voltage-gated channel, Shaw-related subfamily, member 3
 hypothetical protein similar to ankyrin repeat-containing protein AKR1
 hypothetical protein FLJ20533
 hypothetical protein FLJ22643
 solute carrier family 22 (organic cation transporter), member 1-like
 fortheed box Q38
 putative UDP-GalNAc:polypeptide N-acetylglucosaminyltransferase T9
 peptidylprolyl isomerase (cyclophilin)-like 2
 glucagon receptor
 Consensus includes gb:AC004490/DEF=Homo sapiens chromosome 19, cosmid R29381/FEA=CDS/DB_XREF=gj:2995576/UG=Hs.156015 Homo sapiens chromosome 19,
 CGI-138 protein
 SH3-domain protein 5 (ponatin)
 sirutin alien mating type information regulation 2 homolog 5 (S. cerevisiae)
 opiate receptor-like 1
 lipase, hormone-sensitive
 type I transmembrane receptor (seizure-related protein)
 95 kDa retinoblastoma protein binding protein
 BAI1-associated protein 3
 apoptosis inhibitor 5
 Consensus includes gb:AL049314.1/DEF=Homo sapiens mRNA; cDNA DKFZp564C156 (from clone DKFZp564C156). /FEA=mRNA/DB_XREF=gj:4500088/UG=Hs.2785921/AL049314
 phosphorylase kinase, alpha 2 (liver)
 serine protease inhibitor, Kunitz type 1
 ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog, yeast)
 cytochrome P450 retinoid metabolizing protein

MOST-1 protein
 hypothetical protein FLJ13330
 proteasome (prosome, macropain) subunit, alpha type 4
 hypothetical protein FLJ23429
 Consensus includes gb:AL339602.1 /DEF=Homo sapiens mRNA; cDNA DKFp547D148 (from clone DKFp547D148). /FEA=mRNA /DB_XREF=gi:8655670 /UG=Hs.306514 t AL339602
 perlecanin-like material 1
 arrestin 3, retinal (k-arrestin)
 histone acetyltransferase
 KIAA0553 protein
 exonuclease NEF-4p
 hypothetical protein FLJ10979
 Consensus includes gb:AK026803.1 /DEF=Homo sapiens cDNA, FLJ23150 fts, clone LNG09338. /FEA=mRNA /DB_XREF=gi:10439742 /UG=Hs.277178 Homo sapiens cDNA; BC006116
 tubulin-specific chaperone c.
 hypothetical protein FLJ11722
 Consensus includes gb:AF236545 /DEF=Homo sapiens chromosome 17, HC90, HC71AC, HC6 and HC56 genes, complete sequence /FEA=CDS /DB_XREF=gi:10945429 /UG=Hs.326457 ESTs
 afamin
 spindle pole body protein
 cell division cycle 25C
 SRY (sex determining region Y)-box 2
 KIAA1862 protein
 nuclear receptor subfamily 2, group F, member 8
 B-cell CLL/lymphoma 7C
 interleukin 17 receptor
 single Ig IL-1R-related molecule
 KIAA1033 protein
 Consensus includes gb:AA648851 /FEA=EST /DB_XREF=gi:2577179 /DB_XREF=est:ns53g06.s1 /CLONE=IMAGE:1187386 /UG=Hs.326457 ESTs
 PR domain containing 4
 GNAS complex locus
 PDZ-binding kinase; T-cell originated protein kinase
 hypothetical protein FLJ21945
 colony stimulating factor 2 (granulocyte-macrophage)
 Consensus includes gb:AA513509 /FEA=EST /DB_XREF=gi:7151587 /DB_XREF=est:xd6d08.x1 /CLONE=IMAGE:2707023 /UG=Hs.254679 ESTs, Highly similar to Y539_H1 AW513509
 TRIAD3 protein
 Consensus includes gb:A659005 /FEA=EST /DB_XREF=gi:4762575 /DB_XREF=est:u23602.x1 /CLONE=IMAGE:2251698 /UG=Hs.4082 lectin, galactoside-binding, soluble, t A659005
 hypothetical protein
 progesterone receptor membrane component 2
 glutamate receptor, ionotropic, kainate 5
 a disintegrin and metalloproteinase domain 28
 LIM domain binding 1
 mannosidase, alpha, class 1B, member 1
 hypothetical protein FLJ20847
 acyl-Coenzyme A oxidase 1, palmitoyl
 hypothetical protein FLJ13693
 splicing factor, arginine/serine-rich 2
 melanoma antigen, family B, 3
 vacuolar related kinase 2
 lamin B receptor
 Bicoid/D homolog 1 (Drosophila)
 pleomorphic adenoma gene-like 1
 similar to S. cerevisiae RER1
 placental protein 13-like protein
 KIAA0406 gene product
 eductin 3 (gamma)
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.5kD, MWFE)
 RAD52 homolog (S. cerevisiae)
 hypothetical protein FLJ22624
 endothelial cell growth factor 1 (platelet-derived)
 proviral dehydrogenase (lipamide) alpha 1
 nuclear factor I/C (CCAAT-binding transcription factor)
 nucleobindin 1
 secreted and transmembrane 1
 protein phosphatase 2A, regulatory subunit B' (PR 53)
 3'-phosphoadenosine 5'-phosphosulfate synthase 2
 AW299958

hypothetical protein FLJ11467	NM_024863	1.09
retinol dehydrogenase homolog	NM_005771	1.09
nuclear protein	AU357871	1.09
MDM2 box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	NM_005587	1.09
down-regulated in metastasis	AF072718	1.09
high-mobility group (nonhistone chromosomal) protein 1	NM_002128	1.09
claudin 11 (oligodendrocyte transmembrane protein)	NM_005602	1.09
corstatin	AL578583	1.09
KIAA1232 protein	AU16274	1.09
muopin 1	NM_020533	1.09
phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 duncce homolog, Drosophila)	U18088	1.09
putative Rab5-interacting protein	NM_018840	1.09
PCTAIRE protein kinase 2	AU742210	1.09
tropomodulin	AI583383	1.09
Domin syndrome critical region gene 3	NM_006052	1.09
seclite 2 (Drosophila)	BC765616	1.09
KIAA0912 protein	AB020719	1.09
Consensus includes gb:AL049842/DEF=Human DNA sequence from clone 129L7 on chromosome 8q12-13. Contains the gene for a PUTATIVE novel protein. ESTs, an STS, 'AL049842	NM_006926	1.09
surfactant, pulmonary-associated protein A2	AU118882	1.09
endothelin receptor type A	BC165833	1.09
fatty acid desaturase 1	BC001889	1.09
N-ethylmaleimide-sensitive factor attachment protein, gamma	NM_014833	1.09
Consensus includes gb:AL109714.1/DEF=Homo sapiens mRNA (full length insert cDNA clone EUROMIMAGE 327506./FEA=mRNA/DB_XREF=gi:5689826/UG=Hs.195484 Ho	AL109714	1.09
KIAA0618 gene product	BF436315	1.09
Consensus includes gb:BF436315/FEA=EST/DB_XREF=gi:11448630/DB_XREF=est:706605.x1/CLONE=IMAGE:3644888/UG=Hs.296338 ESTs	NM_003916	1.09
adaptor-related protein complex 1, sigma 2 subunit	NM_002977	1.09
sodium channel, voltage-gated, type IX, alpha polypeptide	AA172285	1.09
vitamin D (1,25-dihydroxyvitamin D3) receptor	NM_004554	1.09
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4	NM_002921	1.09
retinal G protein coupled receptor	AW188134	1.09
retinoic acid repressible protein	NM_003717	1.09
neuropeptide FF-amide peptide precursor	NM_018655	1.09
lens epithelial protein	Q26054	1.09
luciferase-1 beta-phosphatase 1	S81545	1.09
endothelin receptor type A	NM_001424	1.09
epithelial membrane protein 2	AA143765	1.09
Consensus includes gb:AA143765/FEA=EST/DB_XREF=gi:1713188/DB_XREF=est:z031e02.s1/CLONE=IMAGE:588506/UG=Hs.71779 Homo sapiens DNA from chromosome	AA143765	1.09
CDI-08 protein	NM_015937	1.09
protease (prosome, macropain) subunit, beta type, 5	BC704146	1.09
translocase of inner mitochondrial membrane 8 homolog B (yeast)	NM_012459	1.09
CD8 antigen, beta polypeptide 1 (p27)	AW296309	1.09
KIAA0193 gene product	NM_014738	1.09
hypothetical protein FLJ2222	AI855411	1.08
transforming growth factor, beta receptor II (70-80kD)	NM_003242	1.08
Consensus includes gb:AU146945/FEA=EST/DB_XREF=gi:11008468/DB_XREF=est:AU146945/CLONE=HEMNB1001944/UG=Hs.296687 Homo sapiens cDNA FLJ12040	AU146945	1.08
aldohydroxy dehydrogenase 5 family, member A1 (succinate-semialdehyde dehydrogenase)	NM_001080	1.08
KIAA1096 protein	NM_001080	1.08
Consensus includes gb:NA48361/FEA=EST/DB_XREF=gi:1189527/DB_XREF=est:yy79g10.s1/CLONE=IMAGE:279810/UG=Hs.15787 Homo sapiens mRNA: cDNA DKFZp5	NA48361	1.08
GTP-D-glucose 4 beta-dehydratase	NM_014305	1.08
ADP-ribosylation factor domain protein 1, 64kD	AF230398	1.08
hypothetical protein MGC3222	W74580	1.08
TRIA03 protein	NM_019011	1.08
actin cross-linking factor	AK023406	1.08
transferrin receptor 2	NM_003227	1.08
Consensus includes gb:AL135960/DEF=Human DNA sequence from clone RP1-18D14 on chromosome 1 Contains genes for SIL (TAL1 (SCL) interrupting locus), TAL1 (T-ce	NM_017697	1.08
hypothetical protein FLJ20171	X96507	1.08
myosin IC	AF201385	1.08
thioredoxin reductase beta	BC476681	1.08
cell division cycle 34	AF041410	1.08
malignancy-associated protein	AF038905	1.08
linker for activation of T cells	AB042823	1.08
RecQ protein-like 5	NM_004593	1.08
splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)		

Cystatin EM	NM_001323	1.08
leucine zipper, putative tumor suppressor 1	AF123656	1.08
Splicing factor, arginine/serine-rich, 48KD	A1343248	1.08
tumor endothelial marker 7 precursor	NM_020405	1.08
acrosomal vesicle protein 1	NM_020115	1.08
hypothetical protein FLJ10314	NM_018055	1.08
hypothetical protein FLJ20272	NM_017735	1.08
hypothetical protein MGC2628	NM_024076	1.08
cubilin (intrinsic factor-cobalamin receptor)	NM_001081	1.08
SKIP for skeletal muscle and kidney enriched inositol phosphatase	NM_016532	1.08
immunoglobulin mu binding protein 2	L14754	1.08
tumor-associated calcium signal transducer 2	J04152	1.08
KIAA0349 protein	A1761518	1.08
sequestrin 1	NM_003900	1.08
KIAA0841 protein	AB020646	1.08
myeloid leukemia factor 1	NM_022443	1.08
poly(A) binding protein, nuclear 1	A130920	1.08
guanylate cyclase activator 2B (uroguanylin)	NM_007102	1.08
cisplatin resistance-associated overexpressed protein	NM_016424	1.08
neuropeptide Y receptor Y2	NM_000910	1.08
mannose-6-phosphate receptor (calnexin dependent)	NM_002355	1.08
KIAA1087 protein	BE646386	1.08
nuclear factor related to kappa B binding protein	NM_006165	1.08
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37KD)	D55672	1.08
MAP1microtubule affinity-regulating kinase like 1	AB049127	1.08
CDK7 cell division cycle 7-like 1 (S. cerevisiae)	NM_017717	1.08
PTD015 protein	NM_003503	1.08
solute carrier family 15 (H ⁺ /peptide transporter), member 2	BC002752	1.08
collectin sub-family member 12	BF223679	1.08
fibroblast growth factor 22	NM_030781	1.08
four and a half LIM domains 1	NM_020637	1.08
DKFZP568J0619 protein	NM_001449	1.08
kinesin heavy chain member 2	BF055496	1.08
glycogenin	BE872663	1.08
topoisomerase (DNA) II alpha (170KD)	AF087842	1.08
chromosome 21 open reading frame 91	AL581834	1.08
testis specific ankyrin-like protein 1	NM_017447	1.08
KIAA0923 protein	NM_018552	1.08
hypothetical protein FLJ20967	AK001710	1.08
collagen, type VI, alpha 2	NM_022071	1.08
CD1D antigen, d polypeptide	AY029208	1.08
ubiquitin-cytochrome c reductase binding protein	NM_001766	1.08
KIAA0176 protein	BC005230	1.08
Consensus includes gb:AW006893 /FEA=EST /DB_XREF=gi:5855731 /DB_XREF=est:wd8606.x1 /CLONE=IMAGE:2506859 /UG=Hs.237946 ESTs, Moderately similar to ALL	A4523921	1.08
imidazole receptor candidate	NM_007184	1.08
hypothetical protein FLJ12118	NM_024537	1.08
hemochromatosis	AF148804	1.08
hypothetical protein FLJ14069	NM_025023	1.08
DKFZP564G196 protein	AL049329	1.08
spolipoprotein A-I	NM_000039	1.08
hypothetical protein FLJ20207	NM_017711	1.08
myosin, light polypeptide 9, regulatory	NM_006097	1.08
radixin	AL137761	1.08
replication factor C (activator 1) 5 (36.5KD)	BC001868	1.08
regulator of G-protein signalling 12	AA687475	1.08
Consensus includes gb:AL046979 /FEA=EST /DB_XREF=gi:5435035 /DB_XREF=est:DKFZp568K0617_s1 /CLONE=DKFZp568K0617 /UG=Hs.8973 tensin	AL046979	1.08
Consensus includes gb:AL574186 /FEA=EST /DB_XREF=gi:12934150 /DB_XREF=est:AL574186 /CLONE=CSDD0039YM09 (3 prime) /UG=Hs.9739 glycerol-3-phosphate dehyd	AL574186	1.08
trichocoid repeat containing 11 (THR-associated protein, 230 kDa subunit)	AF132033	1.08
Consensus includes gb:NT3931 /FEA=EST /DB_XREF=gi:1231216 /DB_XREF=est:z74a07.s1 /CLONE=IMAGE:298260 /UG=Hs.323079 Homo sapiens mRNA; cDNA DKFZp	NT3931	1.08
KIAA0818 protein	NM_015057	1.08
complement component 1, q subcomponent binding protein	M65039	1.08
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 15	NM_001358	1.08

tumor necrosis factor receptor superfamily, member 6	AA164751	1.08
KIAA1449 protein	AI06628	1.08
plasminogen activator, urokinase receptor	AC005525	1.08
TATA box binding protein	NM_003194	1.08
alveoly trypsin-like protease	NM_004262	1.08
zinc finger protein 204	AL021808	1.08
HIV-1 Tat interactive protein, 60 kDa	NM_006388	1.08
hypothetical protein FLJ20199	AK00253	1.08
calcium/calmodulin-dependent serine protein kinase (MAGUK family)	NM_003868	1.08
butyrophilin, subfamily 2, member A1	NM_007049	1.08
cholinergic receptor, muscarinic 4	NM_000741	1.08
butyrophilin, subfamily 2, member A1	U90142	1.08
matrin 3	AA129420	1.08
hypothetical protein FLJ10052	NM_017982	1.08
E4F transcription factor 1	NM_004424	1.08
hypothetical protein FLJ2127	NM_022775	1.08
homoeo box A7	AF026397	1.08
protein kinase, X-linked	NM_005044	1.08
BCL2-like 1	U72386	1.08
stII homolog 1 (Drosophila)	AB011537	1.08
ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)	AB239520	1.08
a disintegrin and metalloproteinase domain 9 (mellin gamma)	NM_003816	1.08
KIAA0038 gene product	NM_014642	1.08
neural killer-tumor recognition sequence	NM_005385	1.08
hypothetical protein MGC11335	NM_030819	1.08
3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (soluble)	BG035985	1.08
paired box gene 9	NM_008194	1.08
hypothetical protein FLJ23380	NM_023076	1.08
Rev/Rax activation domain binding protein-related	U727494	1.08
signal recognition particle 19kD	NM_003135	1.08
KIAA0999 protein	D29805	1.08
UDP-Gal4betaGalNAc beta 1,4- galactosyltransferase, polypeptide 1	NM_021176	1.08
islet-specific glucose-6-phosphatase catalytic subunit-related protein	AB018338	1.08
KIAA0795 protein	NM_001446	1.08
fatty acid binding protein 7, brain	NM_000709	1.08
branched chain keto acid dehydrogenase E1, alpha polypeptide (maple syrup urine disease)	NM_025090	1.08
KIAA1453 protein	NM_162044	1.08
Consensus includes gb-AL162044.1 /DEF=Homo sapiens mRNA: cDNA DKFZp761L0812 (from clone DKFZp761L0812); partial cds. /FEA=mRNA /GEN=DKFZp761L0812 /PF AL162044	AA909035	1.08
collagen, type IV, alpha 2	NM_005738	1.08
ADP-ribosylation factor-like 4	AK022548	1.08
Integrin, alpha 7	NM_030788	1.08
DC-specific transmembrane protein	AB23896	1.08
hypothetical protein FLJ14393	AB020630	1.08
protein phosphatase 1, regulatory (inhibitor) subunit t6B	NM_017802	1.08
hypothetical protein FLJ20397	AA176833	1.08
proline synthetase co-transcribed (bacterial homolog)	Consensus includes gb-AK025343.1 /DEF=Homo sapiens cDNA: FLJ21690 fls, clone COL09538. /FEA=mRNA /DB_XREF=gj:10437841 /UG=Hs.306802 Homo sapiens cDNA: AK025343	1.08
arginyl aminopeptidase (aminopeptidase B)-like 1	NM_018226	1.08
mitochondrial ribosomal protein S10	NM_018141	1.08
protein phosphatase 1, regulatory (inhibitor) subunit 12A	NM_002480	1.08
cathepsin L2	AF070448	1.08
mature T-cell proliferation 1	NM_000600	1.08
ALL1 fused gene from Sq31	NM_014423	1.08
KIAA0368 protein	AB002366	1.08
cholinergic receptor, nicotinic, beta polypeptide 1 (muscle)	NM_000747	1.08
Bart-like homeobox 1	NM_021570	1.08
hypothetical protein FLJ22191	NM_025231	1.08
SP140 nuclear body protein	NM_007237	1.08
ADP-ribosylation factor 6	M57763	1.08
trinucleotide repeat containing 5	BC004423	1.08
thymidine repeat containing 5	AB002330	1.08
KIAA0332 protein	NM_006712	1.08
FAST kinase	AK022131	1.08
thioredoxin-like 2		

brain and reproductive organ-expressed (TNFRSF1A modulator)	NM_004899	1.08
SEC23 protein	NM_007214	1.08
DKFZP586C1820 protein	AV694093	1.08
casein kinase 1, alpha 1	BF310713	1.08
nuclear receptor co-repressor 2	AI498095	1.08
succinate dehydrogenase complex, subunit D, integral membrane protein	NM_003002	1.08
insulin-like growth factor binding protein 7	NM_001553	1.08
Rab geranylgeranyltransferase, beta subunit	U49245	1.08
zinc finger protein 151 (p4Z-67)	NM_003443	1.08
RAP1A, member of RAS oncogene family	NM_002884	1.08
hypothetical protein FLJ10276	NM_018045	1.08
LATS, large tumor suppressor, homolog 1 (Drosophila)	NM_004690	1.08
KIAA0481 gene product	T51252	1.08
hypothetical protein FLJ22679	NM_017698	1.08
nemo-like kinase	NM_016231	1.08
complement component 1, q subcomponent, beta polypeptide	NM_000491	1.08
XPA binding protein 1; putative ATP[GTP]-binding protein	AB044661	1.08
ferrochelatase (protoporphyrin)	AU152635	1.08
placental protein 13	NM_013268	1.08
hemopexin	M35803	1.08
suppressor of variegation 3-9 homolog 1 (Drosophila)	NM_003173	1.08
retinoic acid repressible protein	AF130102	1.08
syntrophin 8	NM_004853	1.08
Friedreich ataxia	NM_000144	1.08
breast cancer 2, early onset	X95152	1.08
glutamate dehydrogenase 1	AJ339331	1.08
hypothetical protein FLJ12584	NM_025139	1.08
pyruvate dehydrogenase (lipoamide) beta	AL117618	1.08
butyrophilin, subfamily 3, member A1	U90552	1.08
STRIN protein	NM_016271	1.08
DKFZP586I2223 protein	AL080214	1.08
neuroblastoma overexpressed gene	NM_002514	1.08
dynlen, axonemal, light polypeptide 4	NM_005740	1.08
CASP8 and FADD-like apoptosis regulator	AF009619	1.08
Consensus includes gb:AA744482 FEA+EST		
testis expressed sequence 14	NM_031272	1.08
UDP glycosyltransferase 1 family, polypeptide A9	NM_021027	1.08
KIAA0489 protein	AB007958	1.08
hypothetical protein FLJ20457	NM_017832	1.08
protein tyrosine phosphatase, receptor type, C	AI809341	1.08
mammian-binding lectin serine protease 1 (C4/C2 activating component of Rar-reactive factor)	D17525	1.08
ceroid-lipofuscinosis, neuronal 2, late infantile (Jensky-Bielschowsky disease)	NM_000391	1.08
HCF-binding transcription factor Zhangfei	AI432137	1.08
erlapin 1	AL044651	1.08
ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1	AV711183	1.08
mitochondrial intermediate peptidase	NM_005932	1.08
clustarin (complement lytic inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	M25915	1.08
hypothetical protein FLJ10751	NM_018205	1.07
flucocidase, alpha-L-1, tissue	NM_000147	1.07
SFRS protein kinase 1	NM_003137	1.07
cholinergic receptor, nicotinic, gamma polypeptide	NM_005199	1.07
actin related protein 2/3 complex, subunit 4 (20 kD)	NM_005718	1.07
cell adhesion molecule-related/down-regulated by oncogenes	NM_016952	1.07
complement C11-like proteinase precursor,	NM_016546	1.07
arylsulfatase D	NM_001669	1.07
hypothetical protein FLJ22160	NM_024585	1.07
mitogen-activated protein kinase 7	NM_002749	1.07
dystonia 1, torsion (autosomal dominant; torsin A)	NM_000113	1.07
glutamate receptor, metabotropic 8	AC000099	1.07
hypothetical protein FLJ12879	NM_024757	1.07
hypothetical protein FLJ23469	NM_024710	1.07
HSPQ40 protein	NM_016098	1.07
protein tyrosine phosphatase type IVA, member 1	AW165960	1.07

cytochrome P450, subfamily 1B (phenobarbital-inducible), polypeptide 6
 Consensus includes gb:AF038194.1 DEF-Homo sapiens clone 23821 mRNA sequence, FEA=mRNA_IDB_XREF=gi:2755914 /UG=Hs.302182 Homo sapiens clone 23821 mf
 S73614 1.07
 Immunoglobulin heavy constant mu
 NM_024505 1.07
 NAADPH oxidase, EF hand calcium-binding domain 5
 BC002775 1.07
 ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)
 BC000467 1.07
 peroxisomal biogenesis factor 16
 NM_000325 1.07
 calnexin-like homeodomain transcription factor 2
 AB000815 1.07
 and hydrocarbon receptor nuclear translocator-like
 NM_000033 1.07
 ATP-binding cassette, sub-family D (ALD), member 1
 Consensus includes gb:A037586/FEA=EST_IDB_XREF=gi:4175955/DB_XREF=est:tc24n12.x1 /CLONE=IMAGE:2065607 /UG=Hs.131067 ESTs, Moderately similar to IBA2_2
 NM_000033 1.07
 hypophosphorylated protein MGIC1314 similar to fucosidase, alpha-L-1, tissue
 AL031320 1.07
 KIAA1731 protein
 AB051518 1.07
 Ras-related associated with diabetes
 NM_004165 1.07
 aminolevulinic, delta-, dehydratase
 NM_000031 1.07
 serine (or cysteine) proteinase inhibitor, clade 1 (neuroserpin), member 2
 NM_008217 1.07
 likely ortholog of mouse variant polydenylation protein CSTF-64
 BF732638 1.07
 sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)
 NM_006309 1.07
 leucine rich repeat (in FLU1) interacting protein 2
 NM_025077 1.07
 hypophosphorylated protein FLJ13949
 NM_002223 1.07
 located 1,4,5-bisphosphate receptor, type 2
 AB040900 1.07
 KIAA1487 protein
 NM_020990 1.07
 creatine kinase, mitochondrial 1 (ubiquitous)
 NM_024730 1.07
 hypophosphorylated protein FLJ22855
 NM_024520 1.07
 hypophosphorylated protein FLJ22555
 BC003622 1.07
 located 1,3,4-bisphosphate 5/6 kinase
 NM_004244 1.07
 CD103 antigen
 NM_015972 1.07
 RNA polymerase 1 16 kDa subunit
 likely homolog of rat kinase D-interacting substance of 220 kDa
 NM_018024 1.07
 hypophosphorylated protein FLJ10204
 SH3-domain binding protein 5 (BTX-associated)
 AK022873 1.07
 chromosome 12 open reading frame 4
 NM_015527 1.07
 DKF2P434P1750 protein
 NM_020374 1.07
 mitochondrial ribosomal protein S12
 NM_013377 1.07
 lipoma HMGIC fusion partner-like 2
 N66833 1.07
 protein receptor overlapping transcript-like 1
 AF135156 1.07
 UDP glycosyltransferase 1 family, polypeptide A8
 NM_015344 1.07
 flamin A, alpha (actin binding protein 280)
 NM_019076 1.07
 hypophosphorylated protein FLB6421
 AI133727 1.07
 hypophosphorylated protein FLJ20294
 TAF15 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 68 kD
 NM_035946 1.07
 exosome component Rpp48
 NM_024791 1.07
 BCL2-antagonist/killer 1
 NM_003487 1.07
 cysteine knot superfamily 1, BMP antagonist 1
 NM_020158 1.07
 glycerophosphate O-acyltransferase
 NM_001188 1.07
 aminopeptidase puromycin sensitive
 NM_013372 1.07
 hypophosphorylated protein FLJ20311
 ADP-ribosylation factor 1
 NM_014236 1.07
 nuclear receptor subfamily 6, group A, member 1
 AJ132583 1.07
 customer protein complex, subunit beta 2 (beta prime)
 NM_017750 1.07
 small proline-rich protein 3
 AF052179 1.07
 peroxisomal biogenesis factor 16
 AF004291 1.07
 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2
 NM_004766 1.07
 electron-transferring-flavoprotein dehydrogenase
 NM_005416 1.07
 vimentin beta (SH3-containing adaptor molecule-1)
 AB016531 1.07
 testis-specific ankyrin motif containing protein
 AF046997 1.07
 EphB1
 AF037261 1.07
 hypophosphorylated protein PRO2800
 NM_019644 1.07
 Consensus includes gb:AL031133 /DEF=Human DNA sequence from clone 281H8 on chromosome 6q25.1-25.3. Contains up to four novel genes, one with similarity to KIAA03
 AF037333 1.07
 hypophosphorylated protein FLJ20154
 AK000161 1.07
 acylcarnitine dehydrogenase 4 family, member A1
 NM_003748 1.07
 Microfilament-associated glycoprotein-2
 U37283 1.07

sequestosome 1
neural precursor cell expressed, developmentally down-regulated 4
chromosome 1 open reading frame 2
tumor suppressing subtransferable candidate 1
Mid-1-related chloride channel 1
lec protein tyrosine kinase
5,10-methylenetetrahydrofolate reductase (NADPH)
G protein-coupled receptor 50
HeLa cyclin-dependent kinase 2 interacting protein
actin related protein 2/3 complex, subunit 2 (34 kD)
A kinase (PRKA) anchor protein 11
intercellular adhesion molecule 1 (ICSA), human rhinovirus receptor
tumor repeat associator with PCTAIRE 2
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9
KIAA0680 gene product
DMC1 dosage suppressor of mecl1 homolog, meiosis-specific homologous recombination (yeast)
ornithine decarboxylase anlyzyme 1
UMP-CMP kinase
24 kDa intrinsic membrane protein
cytochrome P450, subfamily XXVIII (25-hydroxyvitamin D-1-alpha-hydroxylase), polypeptide 1
Consensus includes gb:W48843 /FEA=EST /DB_XREF=gi:1337022 /DB_XREF=est:z44a09.s1 /CLONE=IMAGE:325144 /UG=Hs.285814 sprouty (Drosophila) homolog 4 /FL:W48843
cytochrome P450, subfamily IIO (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6
proprotein convertase subtilisin/kexin type 1 inhibitor
heterogeneous nuclear ribonucleoprotein D-like
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7ND, MNLL)
oxytocin, gamma- (neurophysin II)
gb:NM_017421.1 /DEF=Homo sapiens methyltransferase COQ3 (COQ3), mRNA, /FEA=mRNA /GEN=COQ3 /PROD=methyltransferase COQ3 /DB_XREF=gi:8393176 /FL=gb:NM_017421
ankyrin repeat domain 6
Probe nt1737 (polycystic kidney disease, autosomal recessive, in)
ATP binding protein associated with cell differentiation
serum amyloid A4, constitutive
Homer, neuronal immediate early gene, 3
dynein, axonemal, heavy polypeptide 9
inositol 1,4,5-triphosphate receptor, type 1
Consensus includes gb:AL098814 /DEF=Human DNA sequence from clone RP1-139D8 on chromosome 6p12.1-21.1 Contains genes for GUCA1A (guanylate cyclase activator) AL098814
gb:NM_001222.1 /DEF=Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma (CAMK2G), mRNA, /FEA=mRNA /GEN=CAMK2G /PROD=calcium NM_001222
PRO2000 protein
hypothetical protein FLJ22781
KIAA0701 protein
actinin, alpha 4
active BCR-related gene
solite carrier family 5 (sodium-dependent vitamin transporter), member 6
Consensus includes gb:AU148040 /FEA=EST /DB_XREF=gi:1009561 /DB_XREF=est:AU148040 /CLONE=MAMMA1002428 /UG=Hs.301815 Homo sapiens cDNA FLJ12370 AU148040
Consensus includes gb:U21915.1 /DEF=Human chromosome 5q13.1 clone 5G8 mRNA, /FEA=mRNA /DB_XREF=gi:736412 /UG=Hs.14658 Human chromosome 5q13.1 clone U21915
gb:AF274954.1 /DEF=Homo sapiens PNAS-28 mRNA, complete cds, /FEA=mRNA /PROD=PNAS-28 /DB_XREF=gi:12751070 /UG=Hs.326791 Homo sapiens PNAS-29 mRNA /AF274954
gb:NM_017871.1 /DEF=Homo sapiens hypothetical protein FLJ20542 (FLJ20542), mRNA, /FEA=mRNA /GEN=FLJ20542 /PROD=hypothetical protein FLJ20542 /DB_XREF=gi:NM_017871
dishevelled, dish homolog 3 (Drosophila)
KIAA0301 protein
programmed cell death 2
zinc finger protein 93 homolog (mouse)
activating transcription factor 5
ribosomal protein S13
Ca2+-promoted Ras inactivator
CASP8 associated protein 2
erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked)
nucleoporin 54kD
membrane protein of cholinergic synaptic vesicles
hypothetical protein similar to RNA-binding protein Iark
actin, alpha 2, smooth muscle, aorta
ARPI actin-related protein 1 homolog A, contractin alpha (yeast)
integrin, beta 6
senp9/SUMO-specific protease
glutamate receptor, ionotropic, AMPA 2

1.07	HLA class II region expressed gene KE4
1.07	painimogen
1.07	KIAA0107 gene product
1.07	Consensus includes gb BE780075 /FEA=EST /DB_XREF=gi:10201273 /DB_XREF=est601469954F1 /CLONE=IMAGE:3873157 /UG=Hs.74137 transmembrane trafficking protein BE780075
1.07	HLA-B associated transcript 2
1.07	actn, alpha 1, skeletal muscle
1.07	arachidonate 5-lipoxygenase
1.07	xeroderma pigmentosum, complementation group A
1.07	integrin beta 1 binding protein (melusin) 2
1.07	CAT-50 protein
1.07	Consensus includes gb U08764.1 /DEF=Human EST clone 122887 mariner transposon Hamar1 sequence, /FEA=mRNA /DB_XREF=gi:2231367 /UG=Hs.132967 Human EST -U08764
1.07	ATP-binding cassette, sub-family F (GCN2), member 2
1.07	CLIP-associated protein 2
1.07	protein tyrosine phosphatase type IVA, member 2
1.07	Consensus includes gb AL049280.1 /DEF=Homo sapiens mRNA; cDNA DKF_Zp564K143.1 /FEA=mRNA /DB_XREF=gi:500037 /UG=Hs.155397 /AL049280
1.07	Consensus includes gb AF070536.1 /DEF=Homo sapiens clone 24566 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387893 /UG=Hs.133342 Homo sapiens clone 24566 m8 AF070536
1.07	BCL2/pedionovirus E1B 19kD interacting protein 1
1.07	hypothetical protein FLJ110948
1.07	hypothetical protein MGC3178
1.07	zinc finger protein 10 (KDX 1)
1.07	Consensus includes gb AJ249275 /DEF=Homo sapiens partial MT-HFR gene for methylglutathione reductase /FEA=CDS /DB_XREF=gi:7271187 /UG=Hs.214142 5.10-AJ249275
1.07	dimethylarginine dimethylaminohydrolase 2
1.07	mitogen-activated protein kinase kinase 2
1.07	major intrinsic protein of lens fiber
1.07	spectrin, beta, non-erythrocytic 4
1.07	sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1
1.07	hypothetical protein FLJ22439
1.07	chromosome 1 open reading frame 13
1.07	heterogeneous nuclear ribonucleoprotein L
1.07	reovirin
1.07	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1
1.07	comediectoderm
1.07	hypothetical protein MGC2494
1.07	thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)
1.07	olfactory receptor, family 2, subfamily B, member 8
1.07	surfactant, pulmonary-associated protein C
1.07	KIAA1088 protein
1.07	fucosyltransferase 7 (alpha (1,3) fucosyltransferase)
1.07	bacterial/permeability-increasing protein
1.07	hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase)
1.07	gb BC002877.1 /DEF=Homo sapiens. Similar to hypothetical protein FLJ11585, clone MGC:11258, mRNA, complete cds. /FEA=mRNA /PROT=Similar to hypothetical protein BC002877
1.07	PTPRF Interacting protein, binding protein 1 (liprin beta 1)
1.07	immunoglobulin kappa constant
1.07	KIAA0453 protein
1.07	KIAA0424 protein
1.07	Consensus includes gb AF052103.1 /DEF=Homo sapiens clone 23726 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3360410 /UG=Hs.159157 Homo sapiens clone 23726 m8 AF052103
1.07	phosphatidylethanol-4-phosphate 5-kinase, type I, alpha
1.07	eta variant gene 5 (eta-related molecule)
1.07	myogenic factor 6 (herculin)
1.07	hypothetical protein MGC10731
1.07	RAD17 homolog (S. pombe)
1.07	immunoglobulin lambda locus
1.07	serine (or cysteine) proteinase inhibitor, clade A (alpha 1 antitrypsin, antitrypsin), member 3
1.07	protein tyrosine phosphatase, non-receptor type 13 [APO-1/CD95 (Fas)-associated phosphatase]
1.07	nucleosome assembly protein 1-like 1
1.07	hypothetical protein MGC2863
1.07	gag/gi autoantigen, gp130 subfamily b, macrophage (with transmembrane signal), 1
1.07	tamoxifen-diphosphate tamoxifen transferase 1
1.07	Consensus includes gb AK022331.1 /DEF=Homo sapiens cDNA FLJ12289 /ls, clone MAMMA1001635. /FEA=mRNA /DB_XREF=gi:10433706 /UG=Hs.293925 Homo sapiens AK022331
1.07	kuronidase, alpha-L-
1.07	SWAP-70 protein
1.07	Consensus includes gb A638420 /FEA=EST /DB_XREF=gi:4690654 /DB_XREF=est31g07.x1 /CLONE=IMAGE:2242428 /UG=Hs.25035 chloride intracellular channel 4
1.07	cadh18n, type 2
1.07	A8035305

TIAT1 cytoplasmic granule-associated RNA binding protein	AL046419	1.06
cytochrome P450, subfamily 1D (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6	X18866	1.06
hypothetical protein FLJ20320	NM_017765	1.06
serding nexin 10	NM_010322	1.06
hypothetical protein FLJ23529	AK027182	1.06
ribosomal protein L37	BF216701	1.06
Consensus includes gb:AL023773 /DEF=Human DNA sequence from clone 406C18 on chromosome Xq27.1-27.3. Contains two MAGE (Melanoma Associated Antigen) family	AL023773	1.06
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)	AL570294	1.06
KUAA0759 protein	AB018302	1.06
zinc finger protein 283	NM_005741	1.06
nesca protein	NM_014328	1.06
KUAA0657 protein	AI633851	1.06
odorant-binding protein 2A	NM_014582	1.06
ring finger protein 10	AL578551	1.06
AlpV17 transgene, murine homolog, glomerulosclerosis	NM_002437	1.06
KUAA0441 gene product	NM_014797	1.06
Consensus includes gb:AL137673.1 /DEF=Homo sapiens mRNA: cDNA DKFZp343H0872 (from clone DKFZp343H0872). /FEA=mRNA /DB_XREF=gi:16007841 /UG=Hs.30645	AL137673	1.06
Consensus includes gb:AW972351 /FEA=EST /DB_XREF=gi:8162197 /DB_XREF=est:EST384442 /UG=Hs.293451 ESTs	AW972351	1.06
hypothetical protein FLJ10713	NM_018189	1.06
Consensus includes gb:L43578.1 /DEF=Homo sapiens (clone 115392) mRNA. /FEA=mRNA /DB_XREF=gi:895067 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43578	1.06
mitogen inducible 2	AW469573	1.06
hypothetical protein FLJ22531	NM_024650	1.06
MHC class II transactivator	NM_000246	1.06
zinc finger protein 226	NM_015919	1.06
shuin silent mating type information regulation 2 homolog 6 (S. cerevisiae)	NM_016539	1.06
mitogen-activated protein kinase kinase 5	U71087	1.06
SAR1 protein	AY008268	1.06
TIAT1 cytoplasmic granule-associated RNA binding protein	H85548	1.06
hypothetical protein FLJ12598	NM_024754	1.06
Consensus includes gb:Z95624 /DEF=Human DNA sequence from cosmid U237H1 contains Ras like GTPase and ESTs /FEA=CDS /DB_XREF=gi:2117165 /UG=Hs.248013 P_Z95624	NM_024754	1.06
Consensus includes gb:BF882927 /FEA=EST /DB_XREF=gi:12385739 /DB_XREF=est:602306318F1 /CLONE=IMAGE:4397525 /UG=Hs.324787 soluble carrier family 5 (inositol BF882927	BF882927	1.06
Consensus includes gb:N36926 /FEA=EST /DB_XREF=gi:1158068 /DB_XREF=est:y39e04.s1 /CLONE=IMAGE:273534 /UG=Hs.286361 potassium inwardly-rectifying channel N36926	N36926	1.06
carboxypeptidase M	AV710357	1.06
T-box 8	NM_004608	1.06
G protein-coupled receptor 63	NM_030784	1.06
protein kinase C, theta	L01087	1.06
HSV associated factor	NM_006462	1.06
ribosomal protein L27	NM_000968	1.06
chromosome 17 open reading frame 31	AB018275	1.06
KUAA0008 gene product	NM_014750	1.06
heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1	NM_006041	1.06
hypothetical protein MGC2744	AI083187	1.06
KUAA0943 protein	AL080168	1.06
joined to JAZF1	BF382924	1.06
Fragile X mental retardation, autosomal homolog 2	NM_004860	1.06
cytochrome c	BC052999	1.06
candidate tumor suppressor p33 ING1 homolog	NM_016162	1.06
protein phosphatase 2 (formery 2A), catalytic subunit, alpha isoform	BC000400	1.06
hyaluronan synthase 2	NM_005328	1.06
Smr2-related CBP activator protein	AB002307	1.06
BENE protein	BC003179	1.06
hypothetical protein FLJ20136	NM_017684	1.06
v-naf musculoskeletal fibrosarcoma oncogene homolog (avian)	AF055376	1.06
hypothetical protein FLJ12729	NM_024847	1.06
Consensus includes gb:A050783 /FEA=EST /DB_XREF=gi:3307598 /DB_XREF=est:ow12509.x.1 /CLONE=IMAGE:1637081 /UG=Hs.129064 Homo sapiens chromosome 19, c A050783	NM_024847	1.06
DEADH (Asp-Glu-Ala-Asp-His) box binding protein 1	BC004331	1.06
GAP-associated tyrosine phosphoprotein p67 (Sam68)	N24668	1.06
reproduction 6	NM_006559	1.06
optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia)	D83768	1.06
catechol-O-methyltransferase	NM_025136	1.06
ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with mental retardation)	BC000419	1.06
hypothetical protein MGCA174	NM_018941	1.06
	NM_024319	1.06

DKFZ7564G092 protein	1.06	NM_015601.
sphincterectasia 1 (ultrapontocerebellar ataxia 1, autosomal dominant, ataxin 1)	1.06	NM_000332
hypothetical protein FLJ22871	1.06	NM_024861
KIAA1058 protein	1.06	AL576253
KIAA0101 gene product	1.06	NM_014736
hypothetical protein FLJ10914	1.06	NM_018270
culin 3	1.06	NM_003590
diphtheria toxin resistance protein required for diphtheramide biosynthesis (Saccharomyces)-like 2	1.06	NM_001384
negativoblastic leukemia (translocation) 1	1.06	AB037859
ATP-binding cassette, sub-family B (MDR/TAP), member 1	1.06	AF016535
hypothetical protein FLJ15087	1.06	NM_025104
neural precursor cell expressed, developmentally down-regulated 4-like	1.06	AB007899
PRO1646 protein	1.06	AF116078
Consensus includes gb:BG167370 /FEA=EST /DB_XREF=gi:12874199 /DB_XREF=est:602342795F.1 /CLONE=IMAGE:4453055 /UG=Hs.321717 ESTs, Weakly similar to S22; BG167370	1.06	AF116078
nuclear receptor subfamily 3, group C, member 1	1.06	BC116757
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylglucosaminyltransferase 1 (GalNAc-T1)	1.06	X03348
osteonodulin	1.06	NM_020474
flattin A, alpha (actin binding protein 280)	1.06	A1765819
pryl endopeptidase	1.06	AW051856
nuclear receptor subfamily 4, group A, member 3	1.06	NM_002726
Nef-associated factor 1	1.06	X8984
hypothetical protein MGC5468	1.06	NM_006058
inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase gamma	1.06	A0681536
frizzled homolog 1 (Drosophila)	1.06	AF074382
dustrien (complement lytic inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	1.06	NM_003505
dUTP pyrophosphatase	1.06	A1982754
Consensus includes gb:CA005756 /DEF=Homo sapiens chromosome 19, fosmid 39347 /FEA=CDS /DB_XREF=gi:3688091 /UG=Hs.247754 Homo sapiens chromosome 19, lc AC005756	1.06	U62891
melanoma antigen, family D, 2	1.06	AF126181
DKFZP344A043 protein	1.06	BF195973
mouse protein S4, Y-linked	1.06	NM_007008
ras protein-specific guanine nucleotide-releasing factor 1	1.06	L26584
ras homolog gene family, member F (in filopodia)	1.06	NM_018034
pellico homolog 1 (Drosophila)	1.06	NM_020651
galactokinase 1	1.06	BG474736
hypothetical protein FLJ21034	1.06	AK022242
myoendrium collector sultrase	1.06	NM_017947
hypothetical protein FLJ20315	1.06	NM_017763
aminomethyltransferase (glycine cleavage system, protein T)	1.06	NM_000481
bone morphogenetic protein 3 (osteogenic)	1.06	NM_001201
interleukin 9 receptor	1.06	NM_002186
Consensus includes gb:BG235929 /FEA=EST /DB_XREF=gi:12749778 /DB_XREF=est:nf1b02.x1 /CLONE=IMAGE:4141658 /UG=Hs.28034 ESTs	1.06	BG235929
Consensus includes gb:AK024879.1 /DEF=Homo sapiens cDNA: FLJ21028 fls, clone CAE06812 /FEA=mRNA /DB_XREF=gi:10437019 /UG=Hs.317565 Homo sapiens cDNA: AK024879	1.06	BG235929
Consensus includes gb:AW166925 /FEA=EST /DB_XREF=gi:5398450 /DB_XREF=est:tg8601.x1 /CLONE=IMAGE:2633486 /UG=Hs.289037 Homo sapiens cDNA FLJ14135	1.06	AW166925
karyopherin alpha 1 (importin alpha 5)	1.06	AA652173
KIAA1078 protein	1.06	AF593213
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	1.06	BE222436
chromosome 12, open reading frame 2	1.06	NM_007211
carbohydrate (chondroitin) synthase 1	1.06	NM_014918
hypothetical protein FLJ10914	1.06	NM_020995
Chatter (int. AA532952:mf0511.s1) Homo sapiens cDNA, 3' end /clone=IMAGE:1090172 /clone_end=3' /gb=AA582932 /gi=2360292 /ug=Hs.21349 /len=614	1.06	AA582932
hippocampin-related protein	1.06	NM_006822
CD-binding protein homologous to Saccharomyces cerevisiae SEC4	1.06	AF103529
immunoglobulin kappa constant	1.06	AF103529
Consensus includes gb:AB050195.1 /DEF=Homo sapiens mRNA, partial cds, clone:H2-5, MJD homolog /FEA=mRNA /DB_XREF=gi:11559487 /UG=Hs.302079 Homo sapiens AB050195	1.06	AF103529
Consensus includes gb:576346.1 /DEF=AML1 tAML1 (alternatively spliced, exons 5 and 6) human, mRNA Partial, 284 nt /FEA=mRNA /GEN=AML1 /PROCD=AML1 /DB_XREF S76346	1.06	AF103529
Consensus includes gb:BC005066.1 /DEF=Homo sapiens, clone IMAGE:3347854, mRNA, partial cds /FEA=mRNA /PROCD=Unknown (protein for IMAGE:3347854) /DB_XREF BC005066	1.06	AF103529
early endosome antigen 1, 193kD	1.06	NM_003566
33424 B subtilis dapB, pif, pif genes corresponding to nucleotides 1358-3197 of L38424 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	1.06	L38424
KIAA0561 protein	1.06	AB011133
BAT1-associated protein 3	1.06	NM_003933
hypothetical protein FLJ23305	1.06	NM_025059
Consensus includes gb:R87507 /FEA=EST /DB_XREF=gi:946320 /DB_XREF=est:ym91006.s1 /CLONE=IMAGE:166259 /UG=Hs.12509 Homo sapiens clone 23583 mRNA sec. R87507	1.06	NM_025059
forward box E3	1.06	NM_012186
interferon regulatory factor 4	1.06	D78261

protein phosphatase 2, regulatory subunit B (B56), delta isoform NM_006245 1.05
 UDP-N-acetylglucosamine-6-phosphate 4-epimerase (GALNAc-4-epimerase) AF11514 1.05
 RAB14, member RAS oncogene family AF112206 1.05
 androgen receptor (dihydrotestosterone receptor; testicular feminization; spinal and bulbar muscular atrophy; Kennedy disease) AF182704 1.05
 Consensus includes gb:AL534702 /FEA=EST /DB_XREF=gi:12798195 /DB_XREF=est:AL534702 /CLONE=CSODP066TN17 (3 prime) /UG=Hs.13377 Homo sapiens clone 23c AL534702 1.05
 hypothetical protein FLJ10847 NM_018242 1.05
 polp-87 AF204231 1.05
 uncoupling protein 2 (mitochondrial, proton carrier) U94592 1.05
 Consensus includes gb:AU146580 /FEA=EST /DB_XREF=gi:11008101 /DB_XREF=est:AU146580 /CLONE=HEMBB1000893 /UG=Hs.296677 Homo sapiens cDNA FLJ11956 AU146580 1.05
 hypothetical protein FLJ20334 NM_017779 1.05
 CamKII-like protein kinase NM_020397 1.05
 hypothetical protein FLJ12761 NM_024545 1.05
 butyrophilin, subfamily 2, member A1 AL121936 1.05
 sorting nexin 13 AA908770 1.05
 CGI-82 protein NM_016026 1.05
 immediate early response 3 NM_003897 1.05
 triucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit) BC004354 1.05
 CD8 antigen AW134823 1.05
 Consensus includes gb:BE299495 /FEA=EST /DB_XREF=gi:183243 /DB_XREF=est:60094774T1 /CLONE=IMAGE:2960610 /UG=Hs.218329 hypothetical protein /FL=gb:NM BE299495 1.05
 ATPase inhibitor precursor NM_016311 1.05
 RAI/GDS-like gene AF186779 1.05
 Consensus includes gb:BF185104 /FEA=EST /DB_XREF=gi:11081827 /DB_XREF=est:7n15a07.x1 /CLONE=IMAGE:3584565 /UG=Hs.56729 lymphocyte-specific protein 1 BF185104 1.05
 Cluster incl. A1524138:th09103.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2117789 /clone_end=3 /gb=AL524138 /gi=4438273 /ug=Hs.23381 /len=834 A1524138 1.05
 cAMP responsive element binding protein-like 1 AU117487 1.05
 DKFZP568C0424 protein AL561657 1.05
 Consensus includes gb:AU147830 /FEA=EST /DB_XREF=gi:11009351 /DB_XREF=est:qt40b01.x1 /CLONE=IMAGE:1871401 /UG=Hs.126901 Homo sapiens cDNA FLJ12285 AU147830 1.05
 ankyrin 1, erythrocytic NM_020479 1.05
 hypothetical protein BC025668 1.05
 hypothetical protein FLJ23282 AW001438 1.05
 early development regulator 1 (homolog of polyomaic 1) NM_004426 1.05
 hypothetical protein PRO2610 NM_018532 1.05
 yeast Sec31p homolog NM_014933 1.05
 KIAA0725 protein AW005531 1.05
 Consensus includes gb:A1246730 /FEA=EST /DB_XREF=gi:3842127 /DB_XREF=est:qt40b01.x1 /CLONE=IMAGE:1871401 /UG=Hs.126901 Homo sapiens mRNA full length i A1246730 1.05
 diacylglycerol kinase, delta (130kD) NM_003648 1.05
 E1B-55kDa-associated protein 5 BC004242 1.05
 uncharacterized hematopoietic stem/progenitor cells protein MDS030 NM_018465 1.05
 RAB30, member RAS oncogene family NM_014488 1.05
 prostaglandin O2 receptor (DPR) U31099 1.05
 hypothetical protein FLJ10715 NM_018190 1.05
 KIAA1735 protein AF070621 1.05
 thyroid hormone receptor interactor 11 AF007217 1.05
 synaptosomal-associated protein, 91 kDa homolog (mouse) NM_014841 1.05
 cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4) A1401296 1.05
 RAB32, member RAS oncogene family NM_006634 1.05
 heme-regulated initiation factor 2-alpha kinase AW007368 1.05
 hypothetical protein FLJ20533 BC002748 1.05
 Consensus includes gb:AL137651.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O0213 (from clone DKFZp434O0213); partial cds. /FEA=mrna /GEN=OKFZp434O0213 /F AL137651 1.05
 ecdonucleotide pyrophosphatase/phosphodiesterase 1 AU130282 1.05
 claudin 17 AU130282 1.05
 Consensus includes gb:AK023451.1 /DEF=Homo sapiens cDNA FLJ13389 fts, clone PLACE1001171, moderately similar to MYOTUBULARIN. /FEA=mrna /DB_XREF=gi:104 AK023451 1.05
 hypothetical protein NM_012131 1.05
 protein x 0001 AK025847 1.05
 major histocompatibility complex, class II, DO alpha NM_002119 1.05
 integrin, beta 4 BF305661 1.05
 letal giant larvae homolog 2 (Drosophila) NM_004524 1.05
 organic cationic transporter-like 4 NM_004803 1.05
 Consensus includes gb:BF589528 /FEA=EST /DB_XREF=gi:11681865 /DB_XREF=est:naa0505.x1 /CLONE=IMAGE:3254144 /UG=Hs.106642 ESTs, Weakly similar to hypot BF589528 1.05
 sh3-associated polypeptide, 18kD AW274856 1.05
 pumilio (Drosophila) homolog 2 BE778078 1.05
 KIAA0033 protein NG4760 1.05
 Consensus includes gb:BF967998 /FEA=EST /DB_XREF=gi:12335213 /DB_XREF=est:602269508F1 /CLONE=IMAGE:4357777 /UG=Hs.21851 Homo sapiens cDNA FLJ1290 BF967998 1.05

transducin (beta)-like 1	AW968555	1.05
RCE1 homolog, prenyl protein protease (S. cerevisiae)	NM_005133	1.05
secreted protein of unknown function	AA156251	1.05
KIA00943 protein	AA537226	1.05
Consensus includes gb:AL050364.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G052 (from clone DKFZp564G052). /FEA=mRNA /DB_XREF=gi:4914598 /UG=Hs.3027611	AL050364	1.05
deformed epidermal autoregulatory factor 1 (Drosophila)	AF088892	1.05
melanocortin 2 receptor (adrenocorticotrophic hormone)	NM_005229	1.05
presenilin 1 (Alzheimer disease 3)	NM_007319	1.05
EPNA3	NM_005233	1.05
hypothetical protein FLJ10867	NM_018249	1.05
pyruvate kinase, liver and RBC	NM_000298	1.05
keratin 9 (epidermolytic palmoplantar keratoderm)	NM_000226	1.05
Consensus includes gb:AL050376.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586J101 (from clone DKFZp586J101). /FEA=mRNA /DB_XREF=gi:4914609 /UG=Hs.322645 H	AL050376	1.05
MAP-kinase activating death domain	AB002356	1.05
growth arrest and DNA-damage-inducible, beta	AF078077	1.05
DKFZP568D213 protein	NM_015513	1.05
inverdin	AF039217	1.05
DKFZP566C1819 protein	AL050374	1.05
E7A-like factor 1 (eta domain transcription factor)	AL559590	1.05
retinoid acid- and interferon-inducible protein (58kD)	N47725	1.05
thyroid hormone receptor coactivating protein	NM_006696	1.05
sodium channel, voltage-gated, type V, alpha polypeptide (long (electrocardiographic) QT syndrome 3)	NM_000335	1.05
G-protein coupled receptor 88	NM_022049	1.05
tumor necrosis factor (ligand) superfamily, member 7	NM_001252	1.05
SUMO-1-specific protease	AK001408	1.05
melanocortin 5 receptor	NM_005913	1.05
variable charge, X chromosome	AF167079	1.05
hypothetical protein MGC5347	AK022348	1.05
uncharacterized bone marrow protein BM036	NM_018453	1.05
PR domain containing 2, with ZNF domain	AB610133	1.05
calcium/calmodulin-dependent protein kinase IV	L24959	1.05
AT-binding transcription factor 1	NM_006885	1.05
KIAA1139 protein	AK025282	1.05
hypothetical protein DKFZp762166	AI341234	1.05
serine/threonine kinase 10	AB015718	1.05
kalikrein 6 (neurosin, zyme)	NM_002774	1.05
hepatocellular carcinoma-associated antigen 58	NM_016436	1.05
hypothetical protein FLJ13409	NM_024617	1.05
epiplakin 1	NM_031308	1.05
Consensus includes gb:AB15458 /FEA=EST /DB_XREF=gi:4875938 /DB_XREF=est:wb9911.1 /CLONE=IMAGE:2313837 /UG=Hs.206984 Human cosmid CRI-JC2015 at O1	AI275690	1.05
myeloid cell leukemia sequence 1 (BCL2-related)	AI28395	1.05
KIAA1332 protein	W22890	1.05
tetranitropeptide repeat domain 4	NM_015176	1.05
KIAA0483 protein	NM_016566	1.05
G-protein coupled receptor SALPR; somatostatin and angiotensin-like peptide receptor	AL109703	1.05
Consensus includes gb:AL109703.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 25206. /FEA=mRNA /DB_XREF=gi:5689812 /UG=Hs.25314 Hom	U66879	1.05
BCL2-antagonist of cell death	AF152480	1.05
protocadherin alpha 2	AK001290	1.05
neurexin 2	AB035286	1.05
DKFZP564C186 protein	NM_015658	1.05
ortholog of mouse integral membrane glycoprotein LIG-1	AB050468	1.05
phosphate regulating gene with homologies to endopeptidases on the X chromosome (hypophosphatemia, vitamin D resistant rickets)	U87284	1.05
hypothetical protein FLJ20422	AI660497	1.05
regulator of G-protein signalling 10	W19676	1.05
KIAA0089 protein	AA135522	1.05
KIAA0700 protein	AB014600	1.05
hypothetical protein MGC5139	BC004815	1.05
protein tyrosine phosphatase, non-receptor type 21	X79510	1.05
RB-associated KRAB repressor	NM_021163	1.05
activin A receptor, type II	NM_001616	1.05
H2A histone family, member X	AA760862	1.05
prepronociceptin	NM_006228	1.05

Consensus includes gb:Z21967.1 /DEF=H.sapiens partial cDNA for homologue of mPOU homeobox protein. /FEA=mRNA /PROD=mPOU homeobox protein /DB_XREF=gi:437 Z21967
 abd-interactor 12 (SH3-containing protein) U31089 1.05
 apolipoprotein L AF323540 1.05
 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 5 NM_013252 1.05
 guanylate cyclase 1, soluble, alpha 2 NM_000655 1.05
 phosphorotransferase, muscle U24183 1.05
 hypothetical C2H2 zinc finger protein FLJ22504 AA845577 1.05
 transforming growth factor, alpha NM_003236 1.05
 complement component 4B NM_000592 1.05
 Consensus includes gb:Z37544 /DEF=H.sapiens gene for phospholipase C beta 3, exon 1 /FEA=mRNA /DB_XREF=gi:509185 /UG=Hs.37121 phospholipase C, beta 3 (phospholipase C, beta 3) Z37544 1.05
 YY1 transcription factor A4748649 1.05
 zinc finger protein 287 AU150728 1.05
 SOX29 AF032454 1.05
 gb:BC000443.1 /DEF=Homo sapiens. Similar to pericentriolar material 1, clone MGC-8458. mRNA, complete cds. /FEA=mRNA /PROD=Similar to pericentriolar material 1 /DB_XREF=gi:10984291 /DB_XREF=est:7187g06.x1 /CLONE=IMAGE:3391546 /UG=Hs.130465 ESTs BC000443 1.05
 HBV associated factor BE788439 1.05
 hypothetical protein FLJ20979 NM_024121 1.05
 chromosome 22 open reading frame 3 NM_012285 1.05
 hypothetical protein FLJ13593 NM_024780 1.05
 KIAA0515 protein AK028025 1.05
 methylcrotonyl-Coenzyme A carboxylase 2 (beta) AWA39494 1.05
 ceroid-lipofuscinosis, neuronal 3, juvenile (Beitlen, Spielmeier-Vogt disease) AF015593 1.05
 BUB1 budding inhibited by benzimidazoles 1 homologue (yeast) AF043294 1.05
 KIAA0378 protein AK023621 1.05
 Ras association (RAIGDS/AF-6) domain family 1 NM_007182 1.05
 FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (chondrocyte-derived) BF213279 1.05
 fatty acid desaturase 3 NM_021727 1.05
 PAP-1 binding protein BF223045 1.05
 orosomucoid 2 NM_006608 1.05
 histamine receptor H2 NM_022304 1.05
 retinol dehydrogenase 8 (all-trans) NM_015725 1.05
 ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-like 2 AK001880 1.05
 caudal type homeo box transcription factor 1 NM_001804 1.05
 general transcription factor IIC, polypeptide 2 (beta subunit, 110kD) BC000212 1.05
 argininosuccinate synthetase NM_000050 1.05
 KIAA1638 protein NM_025132 1.05
 DKFZP566B163 protein BC028888 1.05
 ter upstream element (FUSE) binding protein 3 U89127 1.05
 RAN, member RAS oncogene family BF112006 1.05
 hypothetical protein PP581 NM_025207 1.05
 hypothetical protein FLJ10734 NM_018187 1.05
 susceptibility protein NSG-x NM_018349 1.05
 protease, serine, 2 (trypsin 2) AF009664 1.05
 inositol polyphosphate-1-phosphatase NM_002194 1.05
 hypothetical protein FLJ23558 NM_025095 1.05
 nuclear receptor subfamily 1, group D, member 2 N32859 1.05
 heterogeneous nuclear ribonucleoprotein A1 AL588186 1.04
 chromobox homolog 1 (HP1 beta homolog Drosophila) NM_006807 1.04
 dolichyl-phosphate mannosyltransferase polypeptide 3 NM_018973 1.04
 truncated calcium binding protein NM_016175 1.04
 Consensus includes gb:A172139 /FEA=EST /DB_XREF=gi:4334229 /DB_XREF=est:186a10.x1 /CLONE=IMAGE:2148378 /UG=Hs.9683 DnaJ (Hsp40) homolog, subfamily C, A472139 1.04
 opsin 1 (cone pigments), short-wave-sensitive (color blindness, tritan) NM_001708 1.04
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 A1860431 1.04
 TRAF interacting protein NM_005879 1.04
 Consensus includes gb:AK024252.1 /DEF=Homo sapiens cDNA FLJ14190 f1s, clone NT2RP2008534, moderately similar to 5-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC SUBUNIT (isoform 1) AK024252 1.04
 KIAA0644 gene product AV724192 1.04
 hypothetical protein FLJ22968 NM_022842 1.04
 hypothetical protein FLJ12355 NM_024988 1.04
 CASP8 and FADD-like apoptosis regulator U97075 1.04
 T cell receptor delta locus X72501 1.04
 phosphodiesterase 5A, cGMP-specific NM_001083 1.04
 latent transforming growth factor beta binding protein 4 NM_003573 1.04
 shiged-like (fascin homology, sea urchin) (Drosophila) NM_000088 1.04
 Consensus includes gb:BF114815 /FEA=EST /DB_XREF=gi:10984291 /DB_XREF=est:7187g06.x1 /CLONE=IMAGE:3391546 /UG=Hs.130465 ESTs BF114815 1.04

KIAA0974 protein	AW300504	1.04
ketohexokinase (fructokinase)	BC006233	1.04
sorting resin 11	NM_013323	1.04
nitric oxide synthase 2A (inducible, hepatocytes)	L24553	1.04
cryptochrome 2 (photolyase-like)	AB014558	1.04
NCX protein	BC006356	1.04
hypothetical protein FLJ20174	NM_017899	1.04
inhibin, beta A (activin A, activin AB alpha polypeptide)	M13436	1.04
KIAA1032 protein	NM_014956	1.04
regulator of G-protein signalling 20	AF074979	1.04
electron-transferring-flavoprotein dehydrogenase	NM_004453	1.04
PHD finger protein 1	AL050332	1.04
phospholipase D1, phosphatidylcholine-specific	NM_002662	1.04
kinectin 1 (kinesin receptor)	Z22551	1.04
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6	NM_016184	1.04
Consensus includes gb:AL109658 /DEF=Human DNA sequence from clone RP4-776F14 on chromosome 20p12.2-13. Contains the 5' end of the FKBP1A gene for FK506-binding	AL109658	1.04
ubiquitin-activating enzyme E1-like	L13852	1.04
hepatoglobin	NM_005143	1.04
hypothetical protein FLJ10060	NM_017386	1.04
collagen, type XI, alpha 2	U32169	1.04
phosphonucleoside-3-kinase, regulatory subunit 4, p150	BF740111	1.04
lysosomal-associated membrane protein 2	J04183	1.04
gb:NM_014925.1 /DEF=Homo sapiens KIAA1002 protein (KIAA1002). mRNA /FEA=mRNA /GEN=KIAA1002 /PROD=KIAA1002 protein /DB_XREF=gi:7662441 /UG=Hs.10249	NM_014925	1.04
hypothetical protein DKFZp434P2235	AL136880	1.04
hypothetical protein, estradiol-induced	NM_015516	1.04
PRO1914 protein	NM_014106	1.04
Consensus includes gb:U79293.1 /DEF=Human clone 23948 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710274 /UG=Hs.159264 Human clone 23948 mRNA sequence	U79293	1.04
cathepsin E	NM_001910	1.04
aristae-like homeobox 4	NM_021926	1.04
hypothetical protein	AV723686	1.04
ribophorin 1	NM_002950	1.04
collectin sub-family member 10 (C-type lectin)	NM_006438	1.04
pro-melanin-concentrating hormone	NM_002874	1.04
stem cell growth factor; lymphocyte secreted C-type lectin	BC005810	1.04
melastatin 1	NM_002420	1.04
stomatin (EBP72)-like 1	NM_004609	1.04
olfactory receptor, family 10, subfamily J, member 1	NM_012351	1.04
parathyroid	M24398	1.04
hypothetical protein FLJ12150	NM_024736	1.04
karyopherin alpha 5 (importin alpha 6)	NM_002269	1.04
ligase III, DNA, ATP-dependent	NM_002311	1.04
melanoma antigen, family A, 5	AI200443	1.04
solute carrier family 7 (cationic amino acid transporter, y+ system), member 7	NM_003982	1.04
group-specific component (vitamin D binding protein)	NM_000583	1.04
transcription elongation factor B (SII), polypeptide 1-like	NM_003197	1.04
lipoprotein lipase	NM_000237	1.04
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 3	AI760760	1.04
NAD(P)H dehydrogenase, quinone 2	NM_000904	1.04
kinesin family member 5C	NM_004522	1.04
hypothetical protein FLJ14639	AI854627	1.04
Consensus includes gb:AI821657 /FEA=EST /DB_XREF=gi:5440736 /DB_XREF=est:z41809.x5 /CLONE=IMAGE:756184 /UG=Hs.66163 Homo sapiens clone 23636 mRNA	NM_024771	1.04
phosphoenolpyruvate carboxykinase 2 (mitochondrial)	NM_004563	1.04
chemokine (C-C motif) receptor 8	NM_005201	1.04
glutathione S-transferase M5	NM_000851	1.04
lectin, galactoside-binding, soluble, 1 (galectin 1)	NM_002305	1.04
spondyloepiphyseal dysplasia, late	NM_014563	1.04
activated leucocyte cell adhesion molecule	BF242805	1.04
Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA FLJ12241 f1a, clone MAMMA1001274. /FEA=mRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens	AK022303	1.04
SP110 nuclear body protein	NM_004509	1.04
H326	AK076481	1.04
hypothetical protein FLJ12687	NM_024917	1.04
U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 2	NM_005089	1.04

coagulation factor VIII-associated (intronic transcript)	NM_012151	1.04
hepatic leukemia factor	A1810712	1.04
nuclear receptor subfamily 4, group A, member 1	D49728	1.04
CASP8 and FADD-like apoptosis regulator	NM_003879	1.04
ribonuclease 6 precursor	NM_003730	1.04
matrix metalloproteinase 14 (membrane-inserted)	X83535	1.04
hypothetical protein FLJ11753	NM_024659	1.04
protein kinase D2	AF309082	1.04
Ellis van Creveld syndrome	AF216185	1.04
arginine vasopressin receptor 1A	A005066	1.04
fructose-1,6-bisphosphatase 2	NM_003837	1.04
mitochondrial ribosomal protein L48	NM_016055	1.04
Consensus includes gb:NM552205 /FEA=EST /DB_XREF=gi:1198084 /DB_XREF=est:yy44g05.s1 /CLONE=IMAGE:245624 /UG=Hs.20205 hemoglobin, beta pseudogene 1	N55205	1.04
chaperonin containing TCP1, subunit 5 (epsilon)	AF215798	1.04
LIM domain-containing preferred translocation partner in lipoma	AL044018	1.04
PR domain containing 10	NM_020228	1.04
5-hydroxytryptamine (serotonin) receptor 2C	NM_000868	1.04
zinc finger protein 7 (K0X 4, clone HF.16)	NM_003416	1.04
L1 cell adhesion molecule (hydrocephalus, stenosis of aqueduct of Sylvius 1, MASA (mental retardation, aphasia, shuffling gait and adducted thumbs) syndrome, spastic paraplegia)	AF53981	1.04
COL10 protein	NM_015940	1.04
titer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2	AF263617	1.04
Consensus includes gb:AK026161.1 /DEF=Homo sapiens cDNA: FLJ25508 fts, clone HRC11772. /FEA=mRNA /DB_XREF=gi:10438922 /UG=Hs.8859 Homo sapiens, Similar	NM_025073	1.04
hypothetical protein FLJ21168	NM_017526	1.04
leptin receptor gene-related protein	AF308291	1.04
Consensus includes gb:AF308291.1 /DEF=Homo sapiens serologically defined breast cancer antigen NY-BR-46 mRNA, partial cds. /FEA=mRNA /PROD=serologically defined	NM_018448	1.04
TBP-interacting protein	NM_003364	1.04
uridine phosphorylase	AF201941	1.04
DKFZP566O1648 protein	BC001417	1.04
KIAA0788 protein	AF273049	1.04
nuclear protein	U79257	1.04
F-box only protein 21	NM_01390736	1.04
Consensus includes gb:AL380738 /DEF=Human DNA sequence from clone RP11-209J19 on chromosome 13 Contains ESTs, STSs and GSSs. Contains the gene for the GWI	BC008883	1.04
GPAI1P anchor attachment protein 1 homolog (yeast)	NM_005413	1.04
zinc oculis homeobox homolog 3 (Drosophila)	NM_006162	1.04
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	NM_005207	1.04
v-ort sarcoma virus CT10 oncogene homolog (avian)-like	AA725078	1.04
Consensus includes gb:AA725078 /FEA=EST /DB_XREF=gi:2742785 /DB_XREF=est:a08e12.s1 /CLONE=1342222 /UG=Hs.54567 paired box gene 1 /FL=gb:NM_006192.1	NM_024296	1.04
hypothetical protein MGC1203	X17115	1.04
immunoglobulin heavy constant mu	AB014562	1.04
KIAA0682 gene product	O83175	1.04
winged-type MMTV integration site family, member 7A	NM_016611	1.04
potassium inwardly-rectifying channel, subfamily K, member 4	LOC101	1.04
gb:LOC101.1 /DEF=Human IG VH-region gene, complete cds. /FEA=mRNA /GEN=IGH@ /PROD=immunoglobulin heavy chain V-region /DB_XREF=gi:185526 /FL=gb:LOC101.1	NM_000660	1.04
transforming growth factor, beta 1	X96588	1.04
RYK receptor-like tyrosine kinase	NM_024056	1.04
hypothetical protein MGC5576	BC003103	1.04
KIAA0237 gene product	W90796	1.04
zinc finger protein 75 (D8C6)	AV699786	1.04
glycosylphosphatidylinositol specific phospholipase D1	NM_002494	1.04
NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFY1)	NM_002379	1.04
Consensus includes gb:AK025379.1 /DEF=Homo sapiens cDNA: FLJ21726 fts, clone COLF.088. /FEA=mRNA /DB_XREF=gi:10437882 /UG=Hs.318721 Homo sapiens cDNA	AF180527	1.04
docking protein 1, 82KD (downstream of tyrosine kinase 1)	AF054994	1.04
Consensus includes gb:AF054994.1 /DEF=Homo sapiens clone 23832 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005709 /UG=Hs.31290 Homo sapiens clone 23832 mRNA	NM_004587	1.04
ribosome binding protein 1 homolog 180kD (dog)	NM_017868	1.04
hypothetical protein FLJ20535	BF215996	1.04
myosin IB	BC008373	1.04
steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1)	NM_020213	1.04
hypothetical protein from EUROIMAGE 1977056	BC005379	1.04
plasmogen-like	AK022897	1.04
reversion-inducing-cytidine-rich protein with kazal motifs	AF014403	1.04
Consensus includes gb:AK025422.1 /DEF=Homo sapiens cDNA: FLJ21769 fts, clone COLF7354. /FEA=mRNA /DB_XREF=gi:10437930 /UG=Hs.306810 Homo sapiens cDNA	AF050145	1.04
phosphatidic acid phosphatase type 2A		
lunonate 2-sulfatase (Hunter syndrome)		

hypothetical protein LOC55565	NM_017530	1.04
cydin 1	BC50368	1.04
hypothetical protein FLJ13224	NM_024799	1.04
8-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 2	NM_008212	1.04
estrogen receptor 2 (ER beta)	AF051428	1.04
creatine kinase, brain	NM_001823	1.04
HIV-1 rev binding protein 2	AJ50314	1.04
protein phosphatase 2A, regulatory subunit B (PR 53)	BC002545	1.04
phospholipase A2, group IIb	AL158172	1.04
hypothetical protein FKSG32	NM_031307	1.04
KIAA0852 protein	NM_014941	1.04
a disintegrin-like and metalloprotease (repolysin type) with thrombospondin type 1 motif, 13	NM_017587	1.04
ribosomal protein S6 kinase, 90kD, polypeptide 2	BC002363	1.04
period homolog 1 (Drosophila)	AF022991	1.04
TGF beta receptor associated protein-1	NM_004257	1.04
KIAA1041 protein	AK027075	1.04
cathepsin W (lymphopain)	NM_001335	1.04
transmembrane protein 1	U61500	1.04
KIAA0414 protein	NM_014007	1.04
hypothetical protein MDS025	NM_021825	1.04
transcription termination factor, RNA polymerase I	NM_007344	1.04
KIAA0202 protein	BC001328	1.04
protein-serine/threonine kinase gene	Z25431	1.04
hypothetical protein FLJ11142	NM_018338	1.04
nuclear receptor subfamily 2, group F, member 1	AJ951185	1.04
Janus kinase 3 (a protein tyrosine kinase, leukocyte)	U31601	1.04
hypothetical protein	NM_019027	1.04
zinc finger protein 147 (estrogen-responsive finger protein)	NM_005082	1.04
low density lipoprotein receptor (familial hypercholesterolemia)	S70123	1.04
protein phosphatase 1, regulatory (inhibitor) subunit 16B	AB020630	1.04
Rho GTPase activating protein 4	NM_001666	1.04
KIAA0378 protein	Z35645	1.04
similar to ZINC FINGER PROTEIN 85 (ZINC FINGER PROTEIN HPF4) (HTF1) (H. sapiens)	AC006539	1.04
nuclear receptor subfamily 1, group D, member 1	NM_021724	1.04
RAS p21 protein activator (GTPase activating protein) 1	NM_002890	1.04
deoxynucleosyltransferase, terminal	M11722	1.04
protein tyrosine phosphatase, non-receptor type substrate 1	AC004832	1.04
protein kinase C binding protein 1	AB032951	1.04
T-cell leukemia/lymphoma 1B	NM_004918	1.04
thousand and one amino acid protein kinase	NM_004783	1.04
coagulation factor VII-associated (intronic transcript)	AJ218431	1.04
Consensus includes gb:AL035287.1 IDEF-H sapiens gene from PAC 747L4 /FEA=mrna PROD=hypothetical protein /DB_XREF=gi:4200248 /UG=Hs.119254 H sapiens gen	AL035287	1.04
amblyot-sensitive cation channel 2, neuronal	NM_020039	1.04
DKF ZP566D193 protein	BE677493	1.04
37 kDa leucine-rich repeat (LRR) protein	NM_005824	1.04
hypothetical protein FLJ20422	NM_017814	1.04
hypothetical protein FLJ12707	NM_022067	1.04
protein kinase, cAMP-dependent, regulatory, type II, alpha	NM_004157	1.04
centaurin, beta 2	D26069	1.04
Consensus includes gb:AA427737 /FEA=EST /DB_XREF=gi:2111578 /DB_XREF=est:zw30g12.s1 /CLONE=IMAGE:770854 /UG=Hs.110820 ESTs	AA427737	1.04
hypothetical protein FLJ20568	NM_017885	1.04
LIM domain kinase 1	D26309	1.04
Consensus includes gb:AJ350339 /FEA=EST /DB_XREF=gi:4087545 /DB_XREF=est:q33g06.x1 /CLONE=IMAGE:1910362 /UG=Hs.144563 tryptophan hydroxylase (tryptophan	AL350339	1.04
Consensus includes gb:AL567779 /FEA=EST /DB_XREF=gi:12921478 /DB_XREF=est:AL567779 /CLONE=CSODF034YB08 (3 prime) /UG=Hs.84775 Human transposon-like	AL567779	1.04
calmodulin 1 (phosphorylase kinase, delta)	N25325	1.04
protein associated with PRK1	AL136598	1.04
clustrin (complement tyrosin inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	M25915	1.04
kinasin family member 5B	NM_004521	1.04
tumor necrosis factor (ligand) superfamily, member 5 (hyper-IgM syndrome)	NM_000074	1.04
son of sevenless (Drosophila) homolog 2	AJ828605	1.04
solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc) transporter), member 3	BC005136	1.04
KIAA1194 protein	NM_015455	1.04
sef-1 suppressor of lin-12-like (C. elegans)	AJ827770	1.04

NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD) (NADH-coenzyme Q reductase)
 hypothetical protein FLJ14129 NM_005006 1.04
 estrogen receptor binding site associated, antigen, 9 NM_030895 1.04
 Interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD) NM_004215 1.04
 Consensus includes gb:BF221873 /FEA=EST /DB_XREF=gi:11128850 /DB_XREF=est:706003.x1 /CLONE=IMAGE:3578429 /UG=Hs.287586 Homo sapiens cDNA FLJ13848 BF221873 BE049439 1.03
 transcriptional regulator interacting with the PHS-bromodomain 2 NM_014755 1.03
 erythrocyte membrane protein band 4.1-like 2 BF511685 1.03
 ribosomal protein S14 M13934 1.03
 adaptor-related protein complex 4, sigma 1 subunit AF155159 1.03
 huntingtin-associated protein 1 (neuroan 1) AF040723 1.03
 and-3-associated polypeptide, 18kD BF593650 1.03
 adenosine B, fructose-bisphosphate AK026411 1.03
 low density lipoprotein receptor-related protein 6 AF074264 1.03
 hypothetical protein PP1228 AI923458 1.03
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2 NM_005874 1.03
 KIAA1074 protein AL137351 1.03
 transcription factor 2, hepatic; LF-B3; variant hepatic nuclear factor NM_006481 1.03
 penicillin-like 2 (Drosophila) U88870 1.03
 e1a variant gene 2 AV693385 1.03
 mitogen-activated protein kinase 14 NM_001315 1.03
 chondroitin sulfate proteoglycan 6 (bemarcan) NM_001315 1.03
 inositol 1,4,5-trisphosphate 3-kinase A NM_002220 1.03
 hypothetical protein FLJ12987 NM_025170 1.03
 DKFZP564M182 protein AK025446 1.03
 myosin, light polypeptide 2, regulatory, cardiac, slow AF020768 1.03
 interferon regulatory factor 5 NM_002200 1.03
 thrombospondin 4 NM_003248 1.03
 KIAA1072 protein NM_014906 1.03
 cat eye syndrome chromosome region, candidate 1 NM_017424 1.03
 hypothetical protein FLJ10521 BC030373 1.03
 gb:L06668.1 /DEF=Homo sapiens porin (por) mRNA, complete cds and truncated cds. /FEA=mRNA /GEN=por. por /PROD=porin; porin /DB_XREF=gi:190189 /FL=gb:L06668.1 /U88866 1.03
 E74-like factor 4 (ets domain transcription factor) U32845 1.03
 kinesin heavy chain member 2 NM_004520 1.03
 v-akt murine thymoma viral oncogene homolog 2 AA789075 1.03
 eukaryotic translation initiation factor 5A2 NM_020390 1.03
 prostaglandin E receptor 3 (subtype EP3) D38300 1.03
 hypothetical protein FLJ20596 AL512707 1.03
 regulatory factor X, 1 (influences HLA class II expression) NM_002818 1.03
 mucin 4, tracheobronchial NM_004532 1.03
 CLST 11240 protein NM_016438 1.03
 clones Z3887 and Z3775 zinc finger protein NM_021188 1.03
 Consensus includes gb:AL138967 /DEF=Human DNA sequence from clone RP11-149M18 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains the natur W88158 1.03
 KIAA0392 protein AI681120 1.03
 RAN binding protein 2 AU145941 1.03
 CDC14 cell division cycle 14 homolog B (S. cerevisiae) NM_031292 1.03
 hypothetical protein DKFZP434G1415 AF041459 1.03
 CASP8 and FADD-like apoptosis regulator X61006 1.03
 tripartite motif-containing 31 X17013 1.03
 DKFZP434H132 protein AW516789 1.03
 macrophage scavenger receptor 1 AF037351 1.03
 Consensus includes gb:BF793446 /FEA=EST /DB_XREF=gi:12098500 /DB_XREF=est:602254922F1 /CLONE=IMAGE:4347388 /UG=Hs.123122 FSH primary response (LRPF BF793446 D87742 1.03
 KIAA0283 protein M85276 1.03
 granulysin NM_013395 1.03
 protein2008 NM_017647 1.03
 Pab1 homolog 3 (E. coli) NM_008193 1.03
 paired box gene 4 BF055171 1.03
 acyl-Coenzyme A oxidase 3, pristanoyl NM_024818 1.03
 hypothetical protein FLJ23251 AL023753 1.03
 hypothetical protein similar to preferentially expressed antigen of melanoma M11734 1.03
 colony stimulating factor 2 (granulocyte-macrophage) NM_002312 1.03
 Consensus includes gb:AA284903 /FEA=EST /DB_XREF=gi:1927445 /DB_XREF=est:224605.r1 /CLONE=IMAGE:714033 /UG=Hs.306366 Homo sapiens T-cell receptor alpha AA284903
 ligase IV, DNA, ATP-dependent

zinc finger protein 36, C3H type-like 1
 leukotriene b4 receptor (chemokine receptor-like 1)
 nitricase 1
 Rho guanine exchange factor (GEF) 12
 KIAA0097 gene product
 KIAA0077 gene product
 onco-like 1 (Drosophila)
 Consensus includes gb:AT161250 /FEA=EST /DB_XREF=gi:5176917 /DB_XREF=est:w60101.x1 /CLONE=IMAGE:2388489 /UG=Hs.90797 Homo sapiens clone 23820 mRNA : A1761250
 myeloid lymphoid or mixed-lineage leukemia (lithotax homolog, Drosophila); translocated to 2
 CAMP responsive element binding protein 1
 neural retina leucine zipper
 platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)
 pyruvate dehydrogenase kinase, isoenzyme 3
 diacylglycerol kinase, theta (110kD)
 myeloid leukemia factor 1
 KIAA0276 protein
 hypothetical protein MGC13219
 hypothetical protein FLJ14153
 protein tyrosine phosphatase, receptor type, (polypeptide (PTPRF), interacting protein (liprin), alpha 3
 fibroblast growth factor 8 (androgen-induced)
 LntC (bacterial lanthibiotic synthetase component C)-like 1
 KIAA0323 protein
 checkpoint with forkhead and ring finger domains
 serine/threonine kinase 10
 Kell blood group precursor (McLeod phenotype)
 RAS guanyl releasing protein 2 (calcium and DAG-regulated)
 chemokine (C motif) XC receptor 1
 hypothetical protein FLJ20037
 zinc finger protein 65 (HPF4, HTF 1)
 solute carrier family 18 (monocarboxylic acid transporters), member 1
 mitochondrial GTP binding protein
 KIAA0542 gene product
 FSH primary response (LRPR1 homolog, rat) 1
 amelogenin (X chromosome, amelogenesis imperfecta 1)
 beta-carotene 15,15'-dioxygenase
 nuclear factor of activated T-cells 5, tonically-responsive
 tripartite motif-containing 2
 perform 1 (pore forming protein)
 solute carrier family 2 (facilitated glucose transporter), member 1
 mannose receptor, C type 1
 ATP-binding cassette, sub-family C (CFTR/MRP), member 3
 olfactory receptor, family 7, subfamily E, member 38 pseudogene
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG)
 KIAA0217 protein
 hypothetical protein FLJ11585
 Consensus includes gb:AF007132.1 /DEF=Homo sapiens clone 23551 mRNA sequence /FEA=mRNA /DB_XREF=gi:2852606 /UG=Hs.184019 Homo sapiens clone 23551 mRNA
 KIAA0480 gene product
 solute carrier family 22 (organic cation transporter), member 4
 Fc fragment of IgG, high affinity Ia, receptor for (CD64)
 ALL1-fused gene from chromosome 1q
 olfactory receptor, family 7, subfamily A, member 17
 DKFZP434F185 protein
 zinc finger protein 154 (pH2-92)
 Consensus includes gb:AL137145 /DEF=Human DNA sequence from clone RP11-563J2 on chromosome 10 Contains ESTs, STSs, GSSs and a CpG island. Contains a novel
 zinc finger protein 36, C3H type-like 2
 Consensus includes gb:AL137474.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E2423 (from clone DKFZp434E2423) /FEA=mRNA /DB_XREF=gi:6000805 /UG=Hs.24477: AL137474
 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E
 Consensus includes gb:BE045142 /FEA=EST /DB_XREF=gi:8362195 /DB_XREF=est:hn28h02.x1 /CLONE=IMAGE:3023283 /UG=Hs.274907 ESTs
 fibronectin-like 2
 cadherin 6, type 2, X-cadherin (fetal kidney)
 platelet-derived growth factor alpha polypeptide
 ubiquitin specific protease 9, X chromosome (Drosophila fat facets related)

ephrin-A4	NM_005227	1.03
steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1)	AF113128	1.03
hypothetical protein FLJ10661	NM_018172	1.03
zinc finger protein 287	NM_020853	1.03
CGI-01 protein	NM_015935	1.03
YY1 transcription factor	NM_003403	1.03
hypothetical protein FLJ20241	AI689402	1.03
novel protein	NM_017515	1.03
Consensus includes gb:AK021609.1 /DEF=Homo sapiens cDNA FLJ11547 fs, clone HEMBA1002534, /FEA=mRNA /DB =Hs.306605 Homo sapiens c AK021609	BF08_054	1.03
CCAAT/enhancer binding protein (C/EBP), delta	AF080103	1.03
Consensus includes gb:AL080103.1 /DEF=Homo sapiens mRNA; cDNA DKFZ564N2216 (from clone DKFZ564N2216), /FEA=mRNA /DB =Hs.14977	NM_002071	1.03
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide, olfactory type	BG437034	1.03
oncostatin M	AK023924	1.03
deltaX homolog 2 (Drosophila)	NM_002445	1.03
macrophage scavenger receptor 1	X92782	1.03
brifazrin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome)	NM_004584	1.03
RAOS homolog (S. pombe)	AI609861	1.03
hypothetical protein from BCRA2 region	NM_024018	1.03
butyrophilin, subfamily 2, member A3	AI588728	1.03
v-akt sarcoma viral oncogene homolog (evian)	NM_006024	1.03
Tax1 (human T-cell leukemia virus type I) binding protein 1	NM_002697	1.03
POU domain, class 2, transcription factor 1	NM_025128	1.03
MUS81 endonuclease	AW089584	1.03
U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 2	NM_023008	1.03
hypothetical protein FLJ17949	NM_023008	1.03
gb:BC004830.1 /DEF=Homo sapiens. Similar to KIAA0798 gene product, clone MGC:10546, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0798 gene product /DB =Hs.306605	NM_001805	1.03
CCAAT/enhancer binding protein (C/EBP), epsilon	NM_008876	1.03
ubiquitin specific protease 20	NM_012217	1.03
tryptase delta 1	BF971923	1.03
mitogen-activated protein kinase kinase 3	NM_006236	1.03
vesicle-associated membrane protein 1 (synaptobrevin 1)	NM_016830	1.03
hypothetical protein FLJ23389	NM_022763	1.03
dishevelled, dish homolog 1 (Drosophila)	AF006011	1.03
Consensus includes gb:AL133262 /DEF=Human DNA sequence from clone RP1-319D22 on chromosome 6 Contains part of the gene for a CDC5-like protein, the gene for a	N32831	1.03
hypothetical protein FLJ10305	NM_018052	1.03
olfactory receptor, family 8, subfamily A, member 1	NM_003896	1.03
MAOS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	NM_002397	1.03
hypothetical protein FLJ21126	NM_025083	1.03
FK506 binding protein 18 (12.6 kD)	AF322070	1.03
radical fringe homolog (Drosophila)	BF940276	1.03
bone morphogenetic protein 6	NM_001718	1.03
cathepsin S	NM_004079	1.03
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	AF009634	1.03
growth arrest-specific 6	L13720	1.03
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 28	NM_018380	1.03
heat shock transcription factor 1	AI593937	1.03
von Hippel-Lindau binding protein 1	NM_003372	1.03
HSPC034 protein	AV700896	1.03
H2A histone family, member 1	NM_003511	1.03
keratin 10 (epidermolytic hyperkeratosis; keratosis palmaris et plantaris)	M19156	1.03
saroglycan, delta (35kD dystrophin-associated glycoprotein)	NM_000337	1.03
hypothetical protein FLJ10989	AI136679	1.03
discs, large (Drosophila) homolog-associated protein 1	AB000277	1.03
seleconectin W, 1	NM_003009	1.03
breakpoint cluster region	U07000	1.03
hypothetical protein FLJ23329	NM_024958	1.03
PROG314 protein	NM_014131	1.03
kaplin (ectin binding protein)	NM_007059	1.03
hypothetical protein MGC3285	NM_025246	1.03
oxysterol binding protein-like 3	AY008372	1.03
hairy and enhancer of split (Drosophila) homolog 2	NM_019089	1.03

sporadic esophageal dysplasia, late, pseudogene
 gp:AF130075.1 /DEF=Homo sapiens clone FLB9413 PRO2532 mRNA, complete cds. /FEA=mRNA /PROD=PRO2532 /DB_XREF=gi:11493455 /UG=Hs.306961 Homo sapiens
 hypothetical protein FLJ11781
 KOEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3
 KIAA0254 gene product
 calcitonin receptor-like
 homolog of C. elegans smu-1
 mitogen-activated protein kinase kinase 14
 similar to granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1) (H. sapiens)
 deoxyhypusine synthase
 KIAA1010 protein
 UDP-Gal:beta-GlcNAc beta 1,4-galactosyltransferase, polypeptide 5
 helicase-mot
 solute carrier family 16 (monocarboxylic acid transporters), member 3
 interferon 8 signal transducer (gp130, oncostatin M receptor)
 protein kinase C binding protein 1
 hemoglobin, alpha 2
 DnaJ (Hsp40) homolog, subfamily B, member 4
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 6
 hypothetical protein FLJ10826
 KIAA0303 protein
 Consensus includes gp:AF533978.1 /DEF=Homo sapiens clone IMAGE:205688, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507325 /UG=Hs.326722 Homo sapiens clone
 pleckstrin homology, Sec7 and coiled-coil domains 4
 hypothetical protein FLJ12270
 Burkitt lymphoma receptor 1, GTP binding protein
 CD28 antigen (Tp44)
 fibronectin II
 Consensus includes gp:AA319460 /FEA=EST /DB_XREF=gi:1971807 /DB_XREF=est:EST21712 /UG=Hs.204166 Human clone 295, 5cm region surrounding hepatocyte nucle
 cortactin SH3 domain-binding protein
 beta transporter/carrier
 RNA polymerase I transcription factor RRN3
 polymerase (DNA directed), beta
 tropomyosin 1 (alpha)
 ubiquitin specific protease
 Consensus includes gp:AW242916 /FEA=EST /DB_XREF=gi:6576686 /DB_XREF=est:xn2703.x1 /CLONE=IMAGE:2694941 /UG=Hs.71988 Homo sapiens mRNA; cDNA DKF
 Rho GTPase activating protein 8
 major histocompatibility complex, class II, DQ alpha 1
 gene near HD on 4p16.3 with homology to hypothetical S. pombe gene
 tubulin-specific chaperone e
 tryptophan rich basic protein
 hypothetical protein
 sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2
 protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform
 Consensus includes gp:AL137534.1 /DEF=Homo sapiens mRNA; cDNA DKFZp34H1419 (from clone DKFZp34H1419); partial cds. /FEA=mRNA /GEN=DKFZp34H1419 /P
 tachykinin receptor 1
 zinc finger protein 124 (HZF-16)
 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
 lateral giant larva homolog 1 (Drosophila)
 DKFZP364K247 protein
 mature T-cell proliferation 1
 Consensus includes gp:AI696737 /FEA=EST /DB_XREF=gi:4984637 /DB_XREF=est:wc56102.x1 /CLONE=IMAGE:2322651 /UG=Hs.304757 Homo sapiens similar to human b
 regulator of G-protein signalling 14
 tumor antigen SLP-8p
 hypothetical protein FLJ10251
 sek-1 suppressor of lin-12-like (C. elegans)
 hypothetical protein FLJ11323
 glucosyltransferase 1 (glactoside 2-alpha-L-glucosyltransferase, Bombay phenotype included)
 hypothetical protein MGC10986
 tumor necrosis factor receptor superfamily, member 8
 FYTD domain-containing ion transport regulator 3
 RAB27A, member RAS oncogene family
 KIAA0495
 E-1 enzyme

casein kinase 1, gamma 1	AB042563	1.02
transmembrane 4 superfamily member 4	BC001388	1.02
annexin A2	BE908217	1.02
lysardite motif-containing 33	AF220137	1.02
guanine nucleotide binding protein (G protein), gamma 5	NM_005274	1.02
aldehyde oxidase 1	NM_001159	1.02
CUG triplet repeat, RNA binding protein 1	NM_006560	1.02
KUAA028 gene product	NM_014724	1.02
adenosine monophosphate deaminase 1 (isoform M)	NM_000036	1.02
T-cell leukemia translocation altered gene	NM_022171	1.02
TONDU	NM_016267	1.02
PDZ domain protein (Drosophila lnaD-like)	NM_005799	1.02
hypothetical protein FLJ11382	NM_021946	1.02
hypothetical protein FLJ10157	NM_018012	1.02
collagen, type V, alpha 1	A082325	1.02
hypothetical protein FLJ11149	BF340123	1.02
hypothetical protein FLJ20898	NM_024600	1.02
LAP [leucine-rich repeats and PDZ] and no PDZ protein	NM_018214	1.02
actin	AA745954	1.02
v-maf musculosarcoma fibrosarcoma oncogene homolog F (avian)	AL021977	1.02
hypothetical protein FLJ21432	NM_024551	1.02
transcription factor 8 (represses interleukin 2 expression)	NM_030751	1.02
fetal hypothetical protein	NM_013303	1.02
signal-induced proliferation-associated gene 1	NM_006747	1.02
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	NM_005176	1.02
proteasome (prosome, macropain) inhibitor subunit 1 (PI31)	NM_006814	1.02
leucine-rich repeat protein, neuronal 1	NM_002319	1.02
Consensus includes gb:AA457019 /FEA=EST /DB_XREF=gi:2179739 /DB_XREF=est:aa38a06.s1 /CLONE=IMAGE:815506 /UG=Hs.280645 ESTs. Weakly similar to ALU7_H1	NM_002319	1.02
KUAA0182 protein	D80004	1.02
SRB7 suppressor of RNA polymerase B homolog (yeast)	A1685580	1.02
hypothetical protein FLJ22439	AW027523	1.02
testis derived transcript (3 LIM domains)	BC001451	1.02
hypothetical protein FLJ12122	NM_024979	1.02
Consensus includes gb:AA099337 /FEA=EST /DB_XREF=gi:1645257 /DB_XREF=IMAGE:489600 /UG=Hs.15780 ATP-binding cassette, sub-family A	AA099337	1.02
hypothetical protein FLJ21458	NM_024850	1.02
CD47 antigen (Rb-related antigen, integrin-associated signal transducer)	BG230614	1.02
hypothetical protein CG0003	NM_023037	1.02
Consensus includes gb:A188796 /FEA=EST /DB_XREF=gi:5591960 /DB_XREF=est:wk20b07.x1 /CLONE=IMAGE:2412853 /UG=Hs.243901 Homo sapiens cDNA FLJ20738	A188796	1.02
calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma	AA284757	1.02
KUAA0440 protein	AB007800	1.02
gb:L23519.1 /DEF=Human Ig rearranged gamma-chain, V-DK4-JH4b, complete cds. /FEA=mrna /DB_XREF=gi:388020 /FL=gb:L23519.1	L23519	1.02
hypothetical protein	BG223334	1.02
immunoglobulin kappa variable 1OR2-108	X51887	1.02
protein kinase, cAMP-dependent, catalytic, beta	AA130247	1.02
thrombospondin 1	A1812030	1.02
malate dehydrogenase 2, NAD (mitochondrial)	AL520774	1.02
lipase, endothelial	NM_008033	1.02
melanocortin 3 receptor	NM_019888	1.02
xylosylprotein beta1.4-galactosyltransferase, polypeptide 7 (galactosyltransferase I)	AK022586	1.02
hypothetical protein FLJ23375	NM_024956	1.02
interleukin-1 receptor-like 1	AB012701	1.02
chromosome 1 open reading frame 21	NM_008086	1.02
phosphatidylcholine	AF031463	1.02
acute carrier family 16 (monocarboxylic acid transporters), member 3	AA853175	1.02
HERV-H LTR-associating 3	NM_007071	1.02
killer cell lectin-like receptor subfamily G, member 1	AF081875	1.02
guanylate cyclase 1, soluble, beta 2	NM_004129	1.02
integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide)	J03925	1.02
casepase recruitment domain 4	NM_006092	1.02
ornithine aminotransferase (gyrate atrophy)	NM_000274	1.02
arachidonate lipooxygenase 3	NM_021628	1.02
zinc finger protein 278	A1807017	1.02
hypothetical protein FLJ12438	NM_021833	1.02

113

KIAA0238 gene product NM_014785 1.02
pyruvate dehydrogenase kinase, isoenzyme 3 NM_005391 1.02
NS1-associated protein 1 AF037448 1.02
Consensus includes gb:AK022094.1 /DEF=Homo sapiens cDNA FLJ12032 fls, clone HEMBB1001880. /FEA=mRNA /DB_XREF=gi:10433415 /UG=Hs.151194 Homo sapiens c AK022094 1.02
IFP38 AF354444 1.02
KIAA1040 protein NM_000610 1.02
CD44 antigen (homolog function and Indian blood group system) NM_000610 1.02
PTD015 protein BC002752 1.02
albumin M12523 1.02
hypothetical protein FLJ20515 AA134926 1.02
actin binding LIM protein 1 BC002448 1.02
Rho GTPase activating protein 8 Z83838 1.02
low density lipoprotein receptor-related protein 8, apolipoprotein e receptor NM_017522 1.02
splicing factor, arginine/serine-rich 3 BC000914 1.02
collagen, type IV, alpha 2 X05610 1.02
transient receptor potential channel 7 AI051254 1.02
hypothetical protein DKFZP434B044 Consensus includes gb:AL359557.1 /DEF=Homo sapiens mRNA; cDNA DKFZP762O1415 (from clone DKFZP762O1415). /FEA=mRNA /DB_XREF=gi:8655612 /UG=Hs.30850 AL359557 1.02
TBP-interacting protein AL136881 1.02
chromosome 2 open reading frame 6 AL136810 1.02
cyclin A1 NM_018221 1.02
mesenchymal stem cell protein DSCD28 NM_003914 1.02
5-hydroxytryptamine (serotonin) receptor 1D NM_016646 1.02
coagulation factor XII (Hageman factor) NM_000864 1.02
acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain NM_000505 1.02
non-metastatic cells 3, protein expressed in NM_000016 1.02
guanine nucleotide binding protein (G protein), beta polypeptide 1-like NM_002513 1.02
mannosidase, alpha, class 1A, member 2 NM_022446 1.02
DKFZP566C243 protein BE543064 1.02
HSPC041 protein AK021433 1.02
U3 snRNP-associated 55-kDa protein NM_016099 1.02
lung type-I cell membrane-associated glycoprotein NM_004704 1.02
uroplakin 2 NM_013317 1.02
estrogen-related receptor beta NM_008760 1.02
Rho guanine nucleotide exchange factor (GEF) 7 NM_004452 1.02
E3 ubiquitin ligase SMURF1 AA778936 1.02
protein phosphatase methyltransferase-1 AIB54869 1.02
KIAA0377 gene product AL040538 1.02
collagen, type IV, alpha 3 (Goodpasture antigen) NM_014659 1.02
hypothetical protein FLJ10618 U02520 1.02
chromosome 15 open reading frame 2 AL049246 1.02
chimerin (chimaerin) 2 NM_018958 1.02
mitochondrial ribosomal protein L2 NM_004087 1.02
bubby homolog (mouse) NM_015850 1.02
paired box gene 2 NM_003320 1.02
CGI-32 protein AW769732 1.02
TATA box binding protein (TBP)-associated factor, RNA polymerase I, C, 110kD NM_015960 1.02
Janus kinase 2 (a protein tyrosine kinase) AW015313 1.02
KIAA0770 protein AF001362 1.02
asparnylglycosaminidase AA812224 1.02
paired box gene 7 M64073 1.02
D-amino-acid oxidase NM_002584 1.02
annexin A2 NM_001617 1.02
gb:BC002811.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ20378, clone MGC:3513, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ20378 1.02
calbindin 1, (28kD) BC001388 1.02
oligodendrocyte lineage transcription factor 2 AF049895 1.02
retinoic acid receptor responder (tazarotene induced) 1 AIB70776 1.02
chromosome 6 open reading frame 34 NM_014320 1.02
inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase gamma AF091453 1.02
complement component 4A K02403 1.02
KIAA0814 gene product AK027138 1.02
uroporphomogen decarboxylase M14016 1.02
DKFZP434B168 protein AL133049 1.02

BTG family, member 4
 KIA0461 protein
 thoredoxin interacting protein
 CGI-115 protein
 v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)
 pancreatitis-associated protein
 hypothetical protein FLJ10079
 hypothetical protein
 core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related
 hypothetical protein MGC2603
 immunoglobulin superfamily, member 1
 plexin B1
 optic atrophy 1 (autosomal dominant)
 ubiquinol-cytochrome c reductase binding protein
 KIA0578 protein
 hypothetical protein FLJ20306
 Consensus includes gb:AL049369.1 /DEF=Homo sapiens mRNA; cDNA DKF7p586D0518 (from clone DKF7p586D0518). /FEA=mRNA /DB_XREF=gi:4500160 /UG=Hs.25072
 tripartite motif-containing 14
 cadherin 13, H-cadherin (heart)
 melanoma differentiation associated protein-5
 calcium channel, voltage-dependent, beta 1 subunit
 xO10 protein
 histone deacetylase 7A
 hypothetical protein FLJ14320
 chloride channel 2
 COX10 homolog, cytochrome c oxidase assembly protein, heme A; lamesyltransferase (yeast)
 KIAA1816 protein
 putative L-type neutral amino acid transporter
 Consensus includes gb:AA477655 /FEA=EST /DB_XREF=gi:2206289 /DB_XREF=est:zu37h09.s1 /CLONE=IMAGE:740225 /UG=Hs.181307 H3 histone, family 3A
 gamma-aminobutyric acid (GABA) A receptor, gamma 3
 RNA helicase-related protein
 bridging integrator 1
 zinc finger protein 187
 G protein-coupled receptor 85
 Consensus includes gb:AL136967 /DEF=Human DNA sequence from clone RP1.149M18 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains the natur
 small inducible cytokine subfamily A (Cys-Cys), member 25
 KIA0429 gene product
 gamma-glutamyltransferase 1
 histamine N-methyltransferase
 Consensus includes gb:AA436973 /FEA=EST /DB_XREF=gi:2179693 /DB_XREF=est:aa90g11.s1 /CLONE=IMAGE:838628 /UG=Hs.74861 activated RNA polymerase II transcr
 hypothetical protein LOC55585
 DKFZP434A062 protein
 Consensus includes gb:AF007146.1 /DEF=Homo sapiens clone 23686 and 23885 mRNA sequences. /FEA=mRNA /DB_XREF=gi:2852624 /UG=Hs.122075 Homo sapiens chr
 Toll-interacting protein
 hypothetical protein FLJ20489
 putative receptor P2X, ligand-gated ion channel, 5
 solute carrier family 26, member 6
 Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fls, clone COL09829. /FEA=mRNA /DB_XREF=gi:10437851 /UG=Hs.306804 Homo sapiens cDNA: AK025352
 hypothetical protein ET
 retinal degeneration B beta
 calpain 1, (mu1) large subunit
 chorionic somatomammotropin hormone-like 1
 ADP-ribosylation factor domain protein 1, 64kD
 transmembrane 4 superfamily member 3
 formin-binding protein 17
 hypothetical protein FLJ23441
 ubiquitin-conjugating enzyme E2L 3
 complement component 6
 erythropoietin receptor
 small nuclear ribonucleoprotein polypeptide N
 CGI-41 protein
 Consensus includes gb:AK021418.1 /DEF=Homo sapiens cDNA FLJ11356 fls, clone HEMBA1000150, highly similar to Homo sapiens pulative RNA helicase mRNA. /FEA=mr

melan-A	U06654	1.01
microspherule protein 1	NM_005337	1.01
Rho-specific guanine nucleotide exchange factor p114	AB011093	1.01
proline-rich G1a (G-carboxyglutamic acid) polypeptide 1	NM_000950	1.01
KIAA0807 protein	AB047005	1.01
polysialin channel, subfamily K, member 1 (TWIK-1)	NM_002245	1.01
O-linked mannose beta 1,2-N-acetylglucosaminyltransferase	NM_017739	1.01
cell cycle progression 2 protein	NM_004749	1.01
ring finger protein (C3H2C3 type) 6	AF203342	1.01
hypothetical protein FLJ23040	NM_025174	1.01
hypothetical protein MG5350	AW612374	1.01
Consensus includes gb:AL049278.1 /DEF=Homo sapiens mRNA; cDNA DKFZp664l153 (from clone DKFZp664l153). /FEA=mRNA /DB_XREF=gi:4500033 /UG=Hs.16074 Hon	AL049278	1.01
GRQ3 oncogene	NM_002090	1.01
SFRS protein kinase 1	AW082913	1.01
paired box gene 8	BC001060	1.01
hypothetical protein FLJ21478	NM_024618	1.01
pregnancy-zone protein	NM_002864	1.01
interleukin 8 receptor, alpha	NM_006334	1.01
nuclear receptor coactivator 2	A040324	1.01
KIAA0179 protein	A481182	1.01
DKFZP547E1010 protein	AW005776	1.01
Consensus includes gb:AW974812 /FEA=EST /DB_XREF=gi:6168015 /DB_XREF=estEST386917 /UG=Hs.291971 ESTs	AW974812	1.01
Bacillus subtilis /REF=X17013 /DEF=B subtilis lys gene for diaminopimelate decarboxylase corresponding to nucleotides 1008-1263 of X17013 /LEN=1108 (-S, -M, -3 represent	X17013	1.01
KIAA0278 protein	D87486	1.01
microtubule-associated protein 7	AW242297	1.01
HSPC156 protein	NM_014178	1.01
TEA domain family member 3	AF142482	1.01
Cluster incl. A452715/ig1:1h06.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:937403 /clone_end=3 /gb=AA527515 /gi=2289584 /ug=Hs.238570 /len=587	AA527515	1.01
smoothelin	AF064238	1.01
transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma)	U85658	1.01
KIAA0157 protein	D63877	1.01
RAN binding protein 3	NM_007321	1.01
serologically defined breast cancer antigen NY-BR-20	AW025529	1.01
death associated transcription factor 1	NM_022105	1.01
adenosine deaminase, RNA-specific, B1 (homolog of rat REDI)	NM_015833	1.01
Consensus includes gb:U50277.1 /DEF=Human breast cancer suppressor element ishmael Upper CP1 mRNA, partial cds. /FEA=mRNA /PROD=suppressor element ishmael	U50277	1.01
CDP-diacylglycerol-inositol 3-phosphatidyltransferase (phosphatidylinositol synthase)	NM_006319	1.01
Consensus includes gb:A1148659 /FEA=EST /DB_XREF=gi:3677128 /DB_XREF=est:qc69c01.x1 /CLONE=IMAGE:1714848 /UG=Hs.296328 ESTs	A1148659	1.01
hypothetical protein FLJ10242	NM_018036	1.01
ornithine carbamoyltransferase	NM_000531	1.01
SELENOPHOSPHATE SYNTHETASE ; Human selenium donor protein	BC000941	1.01
x O99 protein	NM_020234	1.01
dystrophia myotonica-protein kinase	M87313	1.01
zinc finger protein 24 (K0X 17)	BC003566	1.01
ubiquitin-conjugating enzyme E2E-1 (UBC4/5 homolog, yeast)	AL516159	1.01
insulin-like growth factor binding protein 3	BF340228	1.01
retinoic acid induced 15	U50383	1.01
small inducible cytokine subfamily A (Cys-Cys), member 17	NM_002987	1.01
solute carrier family 4 (anion exchanger), member 1, adaptor protein	NM_018158	1.01
hypothetical protein FLJ10482	NM_018107	1.01
inositol polyphosphate-5-phosphatase, 145KD	U53470	1.01
Consensus includes gb:BE531136 /FEA=EST /DB_XREF=gi:9759781 /DB_XREF=est:601278315F1 /CLONE=IMAGE:3610539 /UG=Hs.8982 ESTs, Highly similar to KIAA1139	BE531136	1.01
KIAA0645 gene product	NM_014662	1.01
sialyltransferase 9 (CMP-NeuAc:glactosylceramide alpha-2,3-sialyltransferase; GM3 synthase)	AF119418	1.01
Consensus includes gb:AU145781 /FEA=EST /DB_XREF=gi:11008312 /DB_XREF=est:AU146791 /CLONE=HEMBB1001531 /UG=Hs.287474 Homo sapiens cDNA FLJ12000	AU146791	1.01
lipidoxin	BE56378	1.01
gb:AF249277.1 /DEF=Homo sapiens cervical cancer suppressor-1 mRNA, complete cds. /FEA=mRNA /PROD=cervical cancer suppressor-1 /DB_XREF=gi:10442005 /UG=Hs.	AF249277	1.01
uncharacterized hematopoietic stem/progenitor cells protein MDS032	BC008005	1.01
KIAA0828 protein	BF515124	1.01
amyloid beta (A4) precursor-like protein 2	BC000373	1.01
nuclear DNA-binding protein	NM_006533	1.01
Consensus includes gb:AK021457.1 /DEF=Homo sapiens cDNA FLJ11395 /fs, clone HEMBA1000594. /FEA=mRNA /DB_XREF=gi:10432649 /UG=Hs.306595 Homo sapiens c	AK021457	1.01
choline kinase-like	NM_005198	1.01

pre-B-cell leukemia transcription factor 3	NM_006195	1.01
bridging integrator 1	AF043899	1.01
KIAA0239 protein	NM_015286	1.01
choline acetyltransferase	NM_020985	1.01
phosphatidylinositol-4-phosphate 5-kinase, type I, alpha	U78577	1.01
Fanconi anemia, complementation group A	BG484069	1.01
minor histocompatibility antigen HB-1	NM_021182	1.01
acidic 82 kDa protein mRNA	NM_014597	1.01
hypothetical protein FLJ10782	NM_018216	1.01
G protein-coupled receptor 6	U18549	1.01
membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125)	NM_005699	1.01
hypothetical protein FLJ13055	NM_022737	1.01
cyclin-dependent kinase (CDC2-like) 10	NM_003674	1.01
ecotropic viral integration site 5	AF008915	1.01
prolaserone (prosome, macropain) subunit, alpha type 2	AL523904	1.01
cathepsin B	NM_001908	1.01
topoisomerase (DNA) III alpha	NM_004618	1.01
splicing factor, arginine/serine-rich 2, interacting protein	AW084759	1.01
thromboxane A synthase 1 (platelet, cytochrome P450, subfamily V)	NM_030984	1.01
Sec23-interacting protein p125	AI32962	1.01
fumarylacetoacetate hydrolase (fumarylacetoacetase)	NM_000137	1.01
SP110 nuclear body protein	AF280094	1.01
hypothetical protein FLJ22353	NM_024587	1.01
glucuronidase, beta	NM_000181	1.01
mannan-binding lectin serine protease 1 (C4/C2 activating component of R-reactive factor)	BC000587	1.01
E1A binding protein p300	AI459462	1.01
DKFZP564A122 protein	N24643	1.01
killer cell lectin-like receptor subfamily C, member 3	NM_002261	1.01
gb:NM_016473.1/DEF=Homo sapiens hypothetical protein (LOC51243), mRNA, /FEA=CDS /GEN=LOC51243 /PROD=hypothetical protein /DB_XREF=gi:7706014 /UG=Hs.26	NM_016473	1.01
KIAA0700 protein	BC001205	1.01
hypothetical protein FKSG63	NM_031221	1.01
HSPCO34 protein	AL031427	1.01
KIAA0977 protein	AK002054	1.01
chromosome X open reading frame 12	NM_003492	1.01
gb:NM_001119.2/DEF=Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 1, mRNA, /FEA=mRNA /GEN=ADD1 /PROD=adducin 1 (alpha) isoform a /DB_XREF=gi:77	NM_001119	1.01
DHHC1 protein	NM_016599	1.01
PCF11p homolog	AB020631	1.01
praje 1	NM_022368	1.01
enigma (LIM domain protein)	AI825846	1.01
high-mobility group 20B	N25429	1.01
peter pan homolog (Drosophila)	BC000535	1.01
KIAA0700 protein	AB014600	1.01
enhancer of zeste homolog 2 (Drosophila)	NM_004456	1.01
hypothetical protein FLJ11280	AL561843	1.01
hypothetical protein FLJ11712	NM_024570	1.01
putative N-acetyltransferase Camello 2	NM_016347	1.01
hypothetical protein FLJ10213	NM_018029	1.01
nucleolar cysteine-rich protein	BC004421	1.01
glutamate-ammonia ligase (glutamine synthase)	U08626	1.01
nucleophosmin (nucleolar phosphoprotein B23, numatrin)	AA191576	1.01
aldolase B, fructose-bisphosphate	AV652403	1.01
Consensus includes gb:AF098968.1 /DEF=Homo sapiens familial Mediterranean fever locus region, mRNA sequence, /FEA=mRNA /DB_XREF=gi:4323318 /UG=Hs.113111 H.AFO98968	N39536	1.01
pm5 protein	gb:Z25422.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete CDS, /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405722 /FL=gb:Z25422	1.01
peroxiredoxin 2	L19185	1.01
homogenisate 1,2-dioxygenase (homogenisate oxidase)	AK78172	1.01
Consensus includes gb:AL109696.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 21920, /FEA=mRNA /DB_XREF=gi:5689771 /UG=Hs.185701 Hom	BC000496	1.01
peroxisomal fattyacylated protein	NM_003062	1.01
slit homolog 3 (Drosophila)	NM_007325	1.01
glutamate receptor, ionotropic, AMPA 3	AL022068	1.01
keratin 18	NM_006169	1.01
nicotinamide N-methyltransferase	NM_003043	1.01
solute carrier family 6 (neurotransmitter transporter, laurine), member 6		

KIAA0487 protein
 Consensus includes gb:AA741028 /FEA=EST /DB_XREF=gi:2779620 /DB_XREF=est:ny69c07.s1 /CLONE=IMAGE:1286412 /UG=Hs.256155 ESTs, Moderately similar to ALU1, AA741028
 hypothetical protein FLJ20262
 tumor necrosis factor, alpha-induced protein 1 (endothelial)
 gqgi autoantigen, golgin subfamily a, 1
 kinesin family member 3C
 hypothetical protein PRO2984
 transcription elongation factor 8 (SUI1), polypeptide 3 (110kD, elongin A)
 Consensus includes gb:AK023081.1 /DEF=Homo sapiens cDNA FLJ13019 fis, clone NT2RP3000738, highly similar to Human mRNA for KIAA0140 gene, /FEA=mRNA /DB_XREF=AK023081
 surfactant, pulmonary-associated protein C
 schiste-scul complex-like 1 (Drosophila)
 cytochrome P450, subfamily IIIA (naphthalene oxidase), polypeptide 4
 Consensus includes gb:AA845825 /FEA=EST /DB_XREF=gi:2931965 /DB_XREF=IMAGE:1414703 /UG=Hs.282721 ESTs, Weakly similar to ALU7, H1 AA845825
 solute carrier family 15 (oligopeptide transporter), member 1
 Consensus includes gb:AL134612 /FEA=EST /DB_XREF=gi:6602799 /DB_XREF=est:DKFZp547E086.s1 /CLONE=DKFZp547E086 /UG=Hs.167038 Homo sapiens clone 237, AL134612
 SMA3
 protein inhibitor of activated STAT3
 ilfeguard
 hypothetical protein FLJ11181
 periaxin
 cholinergic receptor, nicotinic, epsilon polypeptide
 neuregulin 1
 cytokine-like nuclear factor n-pac
 G protein-coupled receptor, family C, group 5, member B
 hypothetical protein R33729.1
 bone morphogenetic protein 1
 hypothetical protein FLJ20506
 KIAA0411 gene product
 p47 protein
 procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide I
 myosin, heavy polypeptide 11, smooth muscle
 Eap-related gene member 2
 MLL espin-like fusion
 ret finger protein-like 3 antisense
 spectrin, beta, non-erythrocytic 1
 mitogen-activated protein kinase 8 interacting protein 3
 chromosome 21 open reading frame 2
 hypothetical protein FLJ23282
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide, Y chromosome
 surfactant, pulmonary-associated protein C
 small inducible cytokine subfamily E, member 1 (endothelial monocyte-activating)
 epolipoprotein L, 6
 protein tyrosine phosphatase, receptor type, C
 phospholipase A2, group X
 ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)
 BH-proteadherin (brain-heart)
 mesenchymal stem cell protein DSC43
 p21(CDKN1A)-activated kinase 4
 keratin 18 (focal non-epidermolytic palmoplantar keratoderma)
 anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2)
 hypothetical protein FLJ23322
 caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
 NADPH oxidase 4
 KIAA0295 protein
 keratin 10 (epidermolytic hyperkeratosis, keratosis palmaris et plantaris)
 integrin, alpha 2b (platelet glycoprotein IIb of fibrin complex, antigen CD41B)
 hypothetical protein FLJ20083
 centrosomal protein 1
 progesterone-associated endometrial protein (placental protein 14, pregnancy-associated endometrial alpha-2-globulin, alpha uterine protein)
 myeloid/lymphoid or mixed-lineage leukemia 2
 sigma receptor (SR31747 binding protein 1)
 KIAA1233 protein
 calcium channel, voltage-dependent, beta 3 subunit

mitogen-activated protein kinase 11	U53442	1.01
HSPC157 protein	NM_014179	1.01
hypothetical protein FLJ20598	NM_017895	1.01
hypothetical protein from clones 23549 and 23762	NM_021226	1.01
sperm autoantigenic protein 17	NM_017425	1.01
chemokine (C-C motif) receptor 1	A021071	1.01
suct1-associated neurotrophic factor target 2 (FGFR signalling adaptor)	NM_008653	1.01
hypothetical protein MG24171	NM_024307	1.01
angiotensin receptor 1	NM_000685	1.01
homeo box B7	AW102783	1.01
hypothetical protein FLJ21156	NM_024602	1.01
transient receptor potential channel 2	A1769824	1.01
ubiquitin-cytochrome c reductase binding protein	NM_006294	1.01
ATPase, aminophospholipid transporter-like, Class I, type 8A, member 2	AU148927	1.01
KIAA0884 protein	AK025604	1.01
potassium inwardly-rectifying channel, subfamily J, member 4	U24056	1.01
titin immunoglobulin domain protein (myotilin)	NM_006790	1.01
G protein-coupled receptor 17	NM_008154	1.01
Cytoplasmic FMRP interacting protein 2	Z94154	1.01
KIAA1822 protein	AL161999	1.01
caldesmon 1	AK026910	1.01
ketohexokinase (fructokinase)	A085060	1.01
KIAA0668 protein	NM_000221	1.01
retinal degeneration, slow (retinitis pigmentosa 7)	AK021690	1.01
enigma (LIM domain protein)	NM_000322	1.01
Tcd37 homolog	NM_005451	1.01
Consensus includes gb:U00770.1 /DEF=Human EST clone 251800 mariner transposon Hsmar1 sequence, /FEA=mrna /DB_XREF=gi:2921608 /UG=Hs.42192 Human EST cl U00770	AF123538	1.01
hypothetical protein 869	AF118089	1.01
KIAA0710 gene product	BC000819	1.01
Ellis van Creveld syndrome	NM_014871	1.01
heat shock 105kD	NM_014556	1.01
signal recognition particle 14kD (homologous Alu RNA binding protein)	BG403660	1.01
vesicle-associated membrane protein 5 (myobrevin)	NM_003134	1.01
mitochondrial ribosomal protein S4	NM_006634	1.01
inner centromere protein antigens (135kD, 155kD)	AL138913	1.01
polymerase (DNA directed), lambda	NM_020238	1.01
mannosyl (alpha-1,6)-glycoprotein beta-1,6-N-acetyl-glucosaminyltransferase	NM_013274	1.01
peptidase D	NM_002410	1.01
hypothetical protein P15-2	NM_000285	1.01
DiGeorge syndrome critical region gene 2	AF201942	1.01
Consensus includes gb:AI273812 /FEA=EST /DB_XREF=gi:3896080 /DB_XREF=astqu0807.x1 /CLONE=IMAGE:1964197 /UG=Hs.169252 Human L-myc protein gene, comp AI273812	H42875	1.01
peptidylglycine alpha-amidating monooxygenase	A022882	1.01
Consensus includes gb:AV728268 /FEA=EST /DB_XREF=gi:10437689 /DB_XREF=eat:AV728268 /CLONE=HTCACH05 /UG=Hs.82845 Homo sapiens cDNA: FLJ21930 fls, ck AV728268	U02520	1.01
Consensus includes gb:AK024208.1 /DEF=Homo sapiens cDNA: FLJ21553 fls, clone COL06328, /FEA=mrna /DB_XREF=gi:10437671 /UG=Hs.306786 Homo sapiens cDNA: AK025206	U02287	1.01
collagen, type IV, alpha 3 (Goodpasture antigen)	NM_022448	1.01
selectin P ligand	NM_006308	1.01
hypothetical protein FLJ21817 similar to Rhoip2	NM_014147	1.01
heat shock 27kD protein 3	AL117663	1.01
HSPC047 protein	AB018322	1.01
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	NM_002005	1.01
KIAA0779 protein	BF973178	1.01
feline sarcoma oncogene	SC001214	1.01
G-2 and S-phase expressed 1	L14482	1.01
leucine carboxyl methyltransferase	NM_030925	1.01
POU domain, class 6, transcription factor 1	NM_003955	1.01
hypothetical protein FLJ12577	BC001018	1.00
STAT induced STAT inhibitor 3	NM_017481	1.00
ribosomal protein L39	NM_024695	1.00
ubiquitin 3	X55005	1.00
hypothetical protein FLJ13993	A000419	1.00
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)		
hypothetical protein		

olfactory receptor, family 1, subfamily E, member 1
 cadherin 2, type 1, N-cadherin (neuronal)
 calcium channel, voltage-dependent, alpha 1F subunit
 RAB27B, member RAS oncogene family
 adaptor-related protein complex 3, sigma 2 subunit
 leukocyte membrane antigen
 retinoic acid receptor responder (tazarotene induced) 1
 thyrotropin-releasing hormone
 protocadherin gamma subfamily A, 11
 ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2)
 SKI-like
 adaptor-related protein complex 2, alpha 2 subunit
 hypothetical protein FLJ22809
 UV radiation resistance associated gene
 cannabinoid receptor 1 (brain)
 KIAA0395 protein
 peroxisomal biogenesis factor 7
 luciferase transferase 2 (secretor status included)
 Ewing sarcoma breakpoint region 1
 Consensus includes gbAJ275397 /DEF=Homo sapiens partial (GVH1 gene for immunoglobulin heavy chain V region, case 1, cell Mo V 94 /FEA=CDS /DB_XREF=gi:7573056 /AJ275397
 G protein-coupled receptor
 gb:NM_003871.1 /DEF=Homo sapiens CDC14 (cell division cycle 14, S. cerevisiae) homolog B (CDC14B), mRNA, /FEA=mRNA /GEN=CDC14B /PROD=S, cerevisiae CDC14
 synovial sarcoma, X breakpoint 4
 KIAA1023 protein
 uncoupling 1A
 hypothetical protein FLJ10925
 RAB3A interacting protein (rab33)-like 1
 caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)
 polymerase (DNA directed), theta
 hypothetical protein FLJ11137
 protocadherin alpha 5
 N-acetylase/N-sulfotransferase (heparan glucosaminyl) 1
 aminopeptidase
 deleted in cancer 1; RNA helicase HDB/DICE1
 ribosomal protein L23a-like 1
 T-box, brain, 1
 fetal Alzheimer antigen
 transducin-like enhancer of split 1 (E(spl) homolog, Drosophila)
 hypothetical protein FLJ10248
 methionyl aminopeptidase 2
 actin cross-linking factor
 bromodomain adjacent to zinc finger domain, 1B
 inositol 1,4,5-trisphosphate receptor, type 2
 CCG triplet repeat binding protein 1
 natural killer cell transcript 4
 Consensus includes gbAA129909 /FEA=EST /DB_XREF=gi:1689667 /DB_XREF=est:znr8808.r1 /CLONE=IMAGE:585091 /UG=Hs.145357 ESTs, Moderately similar to ALU7_1
 hypothetical protein FLJ20127
 KIAA0470 gene product
 protocadherin 11
 Consensus includes gb:A1734111 /FEA=EST /DB_XREF=gi:5055224 /DB_XREF=est:ncd8601.y5 /CLONE=IMAGE:771884 /UG=Hs.115870 ESTs
 gb:NM_024304.1 /DEF=Homo sapiens hypothetical protein MGC4859 similar to HSPA8 (MGC4859), mRNA, /FEA=mRNA /GEN=MGC4859 /PROD=hypothetical protein MGC
 Consensus includes gb:AL049307.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564A216 (from clone DKFZp564A216) /FEA=mRNA /DB_XREF=gi:4500076 /UG=Hs.274505 t-AL049307
 synaptosomal-associated protein, 25kD
 Consensus includes gb:AY983138 /FEA=EST /DB_XREF=gi:8152974 /DB_XREF=est:EST375211 /UG=Hs.302040 Homo sapiens DNA sequence from PAC 434O14 on clone AW983138
 Consensus includes gb:AV721987 /FEA=EST /DB_XREF=gi:10824025 /DB_XREF=est:AV721987 /CLONE=HTBAP810 /UG=Hs.184779 Homo sapiens mRNA; cDNA DKFZp5
 Bacillus subtilis /REF=X04603 /DEF=B subtilis thc; thc genes corresponding to nucleotides 1689-2151 of X04603 /LEN=2073 (-5, -M, -3 represent transcript regions 5 prime, X04603
 homeo box A4
 hypothetical protein FLJ11110
 steroid carrier protein 2
 Consensus includes gb:AW268081 /FEA=EST /DB_XREF=gi:6655111 /DB_XREF=est:x929e03.x1 /CLONE=IMAGE:2767612 /UG=Hs.193408 ESTs, Weakly similar to ALUC_1
 bridging integrator 1
 Consensus includes gb:AW268081 /FEA=EST /DB_XREF=gi:6655111 /DB_XREF=est:x929e03.x1 /CLONE=IMAGE:2767612 /UG=Hs.193408 ESTs, Weakly similar to ALUC_1
 dual-specificity lysine-(Y)-phosphorylation regulated kinase 1A
 D86550

DnaJ (Hsp40) homolog, subfamily B, member 12
 adenylyate kinase 3
 fragile X mental retardation 1
 hypothetical protein FLJ20583
 carboxypeptidase A2 (pancreatic)
 hypothetical protein PR02015
 VRK3 for vacuolar related kinase 3
 proliferation-associated ZGA, 38kD
 Consensus includes gb:M20619 /DEF=Human DNA with a hepatitis B virus surface antigen (HBsAg) gene (complete cds) insertion /FEA=CDS /DB_XREF=gi:184331 /UG=Hs.1 M20919
 calcium binding protein P22
 carbohydrate (N-acetylglucosamine-6-O) sulfotransferase 2
 Consensus includes gb:AB041269.1 /DEF=Homo sapiens mRNA for keratin 19, partial cds, isolate:K19-141. /FEA=mRNA, IGEN=mRNA, IGEN=keratin 19 /PROD=keratin 19 /DB_XREF=gi:77:AB041269
 latexin protein
 T cell receptor delta locus
 KIAA0116 protein
 HSPCO34 protein
 pM5 protein
 Consensus includes gb:BE845861 /FEA=EST /DB_XREF=gi:970172 /DB_XREF=est:7677d05.x1 /CLONE=IMAGE:3288489 /UG=Hs.24064 Homo sapiens mRNA: cDNA DKF B6645861 U64094
 peroxisomal biogenesis factor 7
 HIC1-related gene on chromosome 22
 hypothetical protein FLJ21841
 nuclear receptor subfamily 2, group F, member 2
 collagen, type VI, alpha 1
 sorbitol dehydrogenase
 FK506 binding protein 1B (12.8 kD)
 ankylosis, progressive homolog (mouse)
 CGI-98 protein
 potassium voltage-gated channel, shaker-related subfamily, member 4
 chromosome 21 open reading frame 2
 DNA-binding transcriptional activator
 cytochrome P450, subfamily XOA (steroid 21-hydroxylase, congenital adrenal hyperplasia), polypeptide 2
 Consensus includes gb:AK025191.1 /DEF=Homo sapiens cDNA: FLJ21538 fls, clone COL06151. /FEA=mRNA /DB_XREF=gi:10437854 /UG=Hs.306783 Homo sapiens cDNA: AK025191
 sperm associated antigen 8
 sulfotransferase family 4A, member 1
 potassium inwardly-rectifying channel, subfamily J, member 6
 dolichyl-phosphate mannosyltransferase polypeptide 2, regulatory subunit
 eukaryotic translation initiation factor 4E-like 3
 heterogeneous nuclear ribonucleoprotein A1
 dihydrodipyrithinase
 Consensus includes gb:AL139349 /DEF=Human DNA sequence from clone RP11-281P9 on chromosome 20. Contains ESTs, STSs, GSSs and a CpG island. Contains the 3 p:AL139349
 ATPase, Ca++ transporting, type 2C, member 1
 NP0009 protein
 dopamine beta-hydroxylase (dopamine beta-monoxygenase)
 transition protein 2 (during histone to protamine replacement)
 similar to rat myomegalin
 cyclin T1
 hypothetical protein FLJ11040
 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2
 interleukin-1 receptor-associated kinase M
 gamma-tubulin complex protein 2
 Cluster Inc. A1186668 q62a10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:1745468 /gi=3737304 /ug=Hs.21497 /len=512
 PTPRF Interacting protein, binding protein 2 (lpirin beta 2)
 hypothetical protein FLJ22813
 NCK-associated protein 1
 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2

ATP synthase, H ⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1		
KIAA0586 gene product	BC000931	1,24
CGH05 protein	AB011158	1,24
fragile X mental retardation 2	NM_016408	1,24
protein kinase C, delta	AJ001550	1,24
polo (Drosophila)-like kinase	NM_008254	1,24
glycerol kinase	NM_005030	1,24
hypothetical protein FLJ13782	NM_000187	1,24
peroxisomal biogenesis factor 16	NM_024815	1,24
KIAA0268 protein	AA523441	1,24
transmembrane 6 superfamily member 1	AB006824	1,24
protein phosphatase 2, regulatory subunit B (B56), epsilon isoform	NM_023003	1,24
gb:NM_002913.1 /DEF=Homo sapiens replication factor C (activator 1) 1 (145KD) (RFC1), mRNA, /FEA=mRNA /GEN=RFC1 /PROD=replication factor C (activator 1) 1 (145KD)	NM_008248	1,24
KIAA0978 protein	NM_002813	1,24
chromosome 11 open reading frame 5	AL117518	1,24
signal transducer and activator of transcription 1, 91KD	NM_014205	1,24
RNA polymerase I subunit	M87835	1,24
RAE1 RNA export 1 homolog (S. pombe)	AF008442	1,24
developmentally regulated GTP binding protein 2	NM_003610	1,24
hypothetical protein FLJ20373	BF223206	1,24
hyal oxidase-like 2	NM_017792	1,24
NRAS-related gene	BE251211	1,24
interleukin enhancer binding factor 1	NM_007156	1,23
hypothetical protein FLJ12816	NM_004514	1,23
matrix metalloproteinase 15 (membrane-inserted)	NM_022060	1,23
general transcription factor IIIA	NM_002428	1,23
hypothetical protein MGC5378	BE542815	1,23
POD23 homolog 6 (S. cerevisiae)	BC000927	1,23
adenylyl dehydrogenase 3 family, member B2	NM_002874	1,23
splicing factor 3b, subunit 1, 155KD	AA071510	1,23
adenylyl cyclase-associated protein 2	AW003030	1,23
Consensus includes gb:AL034550 /DEF=Human DNA sequence from clone RPS-1184F4 on chromosome 20q11.1-11.23. Contains the 3' end of gene KIAA0978, two genes for AL034550	N90755	1,23
gb:U95006.1 /DEF=Human D9 splice variant A mRNA, complete cds, /FEA=mRNA /PROD=D9 splice variant A /DB_XREF=gi:2071992 /UG=Hs.37616 Human D9 splice variant	U95006	1,23
KIAA0062 protein	C31887	1,23
transforming growth factor, beta receptor II (70-80KD)	D50683	1,23
raslike protein	BF978689	1,23
HGC8.1.1 protein	AB016900	1,23
hypothetical protein	NM_023070	1,23
KIAA1332 protein	AA636114	1,23
LDL induced EC protein	NM_016202	1,23
NS1-binding protein	AB020657	1,23
hypothetical protein RG083M05.2	AC000064	1,23
hypothetical protein MGC3121	NM_024031	1,23
Consensus includes gb:BF001666 /FEA=EST /DB_XREF=gi:10701941 /DB_XREF=est:7g81d12.x1 /CLONE=IMAGE:3313847 /UG=Hs.12460 Homo sapiens clone 23870 mRNA	NM_024031	1,23
dual specificity phosphatase 6	BC001666	1,23
chromosome 21 open reading frame 66	BC005047	1,23
Consensus includes gb:AA928508 /FEA=EST /DB_XREF=gi:3076797 /DB_XREF=est:om17g03.s1 /CLONE=IMAGE:1541332 /UG=Hs.10762 ESTs	NM_013329	1,23
DKFZP564A2418 protein	AA928508	1,23
Ras-GTPase-activating protein SH3-domain-binding protein	AK002064	1,23
solute carrier family 12 (potassium/chloride transporters), member 7	NM_005754	1,23
hypothetical protein FLJ11132	NM_006598	1,23
DR1-associated protein 1 (negative cofactor 2 alpha)	NM_018335	1,23
H2B histone family, member A	NM_006442	1,23
axatrophin	BE271470	1,23
hypothetical protein FLJ10818	BC003404	1,23
karyopherin (importin) beta 1	AB278444	1,23
cargo selection protein (mannose 6 phosphate receptor binding protein)	BC248565	1,23
hypothetical protein MGC14799	AC004381	1,23
nipsnap homolog 1 (C. elegans)	NM_005817	1,23
erythrocyte membrane protein band 4.1-like 3	BC005995	1,23
inhibitor of growth family, member 1	NM_003634	1,23
golgi SNAP receptor complex member 2	NM_012307	1,23
	NM_005537	1,23
	AW148492	1,23

culin 4B	AV694732	1,22
paired box gene 8	X69699	1,22
microtubular-associated protein 1	NM_005926	1,22
peroxisome biogenesis factor 10	BF339133	1,22
DEADH (Asp-Glu-Alu-AspHis) box polypeptide 16	NM_003587	1,22
mitogen-activated protein kinase 9	U35002	1,22
Consensus includes gb:AK024988.1 /DEF=Homo sapiens cDNA: FLJ21335 fls, clone COL02546, /FEA=mRNA /DB_XREF=gi:10437419 /UG=Hs.303001 Homo sapiens cDNA: AK024988		1,22
enhancer of filamentation 1 (cas-like docking; CK-associated substrate related)	U84317	1,22
hypothetical protein FLJ13044	NM_024698	1,22
lymphoblastic leukemia derived sequence 1	BC002796	1,22
WD-repeat protein	BC001284	1,22
hypothetical protein FLJ22824	NM_024608	1,22
nucleolar protein (KKEO repeat)	NM_006392	1,22
high-mobility group (nonhistone chromosomal) protein isoforms I and Y	AF176039	1,22
unit-related transcription factor 1 (acute myeloid leukemia 1; amt1 oncogene)	D43968	1,22
Consensus includes gb:AL120173 /FEA=EST /DB_XREF=gi:5926072 /DB_XREF=est:DKFZp781C117_s1 /CLONE=DKFZp781C117 /UG=Hs.301663 ESTs	AL120173	1,22
hypothetical protein FLJ20727	BE966019	1,22
stratifin	BC000329	1,22
growth hormone inducible transmembrane protein	AF131820	1,22
ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)	M31213	1,22
nucleolar protein family A, member 2 (HACA small nucleolar RNPs)	BC000009	1,22
mitogen-activated protein kinase kinase 1	BE646618	1,22
lectin, galactoside-binding, soluble, 3 (galectin 3)	BC001120	1,22
Consensus includes gb:AI949687 /FEA=EST /DB_XREF=gi:5741997 DB_XREF=est:wt13g04.x1 /CLONE=IMAGE:2471190 /UG=Hs.173638 transcription factor 7-like 2 (T-cell)	AW006750	1,22
hypothetical protein FLJ20059	NM_005652	1,22
telomeric repeat binding factor 2	NM_005873	1,22
regulator of G-protein signalling 19	AV748469	1,22
SEC14-like 1 (S. cerevisiae)	NM_005930	1,22
meningioma expressed antigen 6 (coiled-coil proline-rich)	NM_004716	1,22
proprotein convertase subtilisin/kexin type 7	NM_004716	1,22
anticoagulant protein	AW055008	1,22
H44 antigen	X91103	1,22
FGF receptor activating protein 1	AL049281	1,22
lysozyme homolog	NM_020426	1,22
tail-like receptor 7	NM_016562	1,22
dynactin intermediate chain 2	NM_023036	1,22
viral simian leukemia viral oncogene homolog B (ras related; GTP binding protein)	NM_002881	1,22
myotubularin related protein 3	AF233438	1,22
Consensus includes gb:AF184174 /DEF=Homo sapiens somatostatin receptor 2 (SSTR2) gene, complete cds, alternatively spliced /FEA=CDS_2 /DB_XREF=gi:7229402 /UG=Hs.184174		1,22
Consensus includes gb:AL034396 /DEF=Human DNA sequence from clone 1158812 on chromosome Xp11.21-11.4 Contains the ZXDA gene for X-linked duplicated Zinc finger	AL034396	1,22
RNA helicase-related protein	AF078844	1,22
WD repeat domain 8	NM_017818	1,21
fibronectin 1	NM_024874	1,21
heparan sulfate 2-O-sulfotransferase	BC005858	1,21
hypothetical protein FLJ10468	AK023260	1,21
HERV-H LTR-associated 2	BC001651	1,21
hypothetical protein FLJ11413	NM_007072	1,21
solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11	NM_024554	1,21
KIAA0210 gene product	NM_003562	1,21
JMS protein	A030084	1,21
hypothetical protein DKFZp434E2216	BC000484	1,21
GTT1 protein	BC001043	1,21
RNA guanylyltransferase and 5'-phosphatase	NM_020151	1,21
thymidine synthetase	AB012142	1,21
Consensus includes gb:AL133099.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434B1521 (from clone DKFZp434B1521), /FEA=mRNA /DB_XREF=gi:6453555 /UG=Hs.27231	NM_001071	1,21
Consensus includes gb:BE669922 /FEA=EST /DB_XREF=gi:10318698 /DB_XREF=est:601446588F1 /CLONE=IMAGE:3650432 /UG=Hs.161307 H3 histone, family 3A	BE669922	1,21
nuclear receptor subfamily 0, group B, member 1	NM_000475	1,21
stanniocalcin 2	BC000658	1,21
heat shock transcription factor 2	BC005329	1,21
topoisomerase (DNA) III alpha	AI876092	1,21
Consensus includes gb:AK024175.1 /DEF=Homo sapiens cDNA FLJ14113 fls, clone MAMMA1001715, /FEA=mRNA /DB_XREF=gi:10436491 /UG=Hs.323409 Homo sapiens	AK024175	1,21
heme-regulated initiation factor 2-alpha kinase	NM_014413	1,21

breast cancer metastasis-suppressor 1	AL050008	1,21
hypothetical protein RG083M05.2	AK026142	1,21
mitogen-activated protein kinase 7	U29725	1,21
G protein-coupled receptor 3b	AL137000	1,21
BTB and CNC homology 1, basic leucine zipper transcription factor 2	NM_021813	1,21
CD27-binding (Siva) protein	AW024335	1,21
uncharacterized hematopoietic stem/progenitor cells protein MDS028	NM_018463	1,21
NAD kinase	BE674658	1,21
PFTAIRS protein kinase 1	AF119833	1,21
melanoma-associated antigen recognised by cytotoxic T lymphocytes	NM_006428	1,21
1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)	U56418	1,21
KIAA1035 protein	NM_015239	1,21
F-box only protein 7	NM_012179	1,21
hypothetical protein DKFZp782A227	BC003163	1,21
Sle20-related serine/threonine kinase	NM_014720	1,21
KIAA0794 protein	BG339560	1,21
mitochondrial ribosomal protein L16	NM_017840	1,21
death effector domain-containing	AK022531	1,21
KIAA1449 protein	AB040882	1,21
Consensus includes gb:AI291720 /FEA=EST /DB_XREF=gi:3934494 /DB_XREF=est:cm80.c06.x1 /CLONE=IMAGE:1855050 /UG=Hs.262642 ESTs	AZ91720	1,21
hypothetical protein FLJ22551	NM_024708	1,21
heterogeneous nuclear ribonucleoprotein A1	AI144007	1,21
solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11	AF070548	1,21
hypothetical protein FLJ20258	AF282167	1,21
neurocalcin delta	AF251061	1,21
GATA binding protein 3	AJ796169	1,21
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)	AF100540	1,21
sterol regulatory element binding transcription factor 2	BE513151	1,21
oxoreductase UCPA	NM_020139	1,21
hypothetical protein FLJ12171	NM_024619	1,21
a disintegrin and metalloproteinase domain 19 (metrin beta)	Y13786	1,21
ATP-binding cassette, sub-family G (WHITE), member 5 (sterolin 1)	NM_022436	1,21
zinc finger protein 211	NM_066385	1,21
calcium channel, voltage-dependent, P/Q type, alpha 1A subunit	NM_023035	1,21
hypothetical protein	AC002550	1,21
KIAA1042 protein	A1633774	1,21
brain and reproductive organ-expressed (TNFRSF1A modulator)	U19178	1,21
cyclin G1	BC000106	1,21
arylalkylamine N-acetyltransferase	NM_001088	1,21
KIAA0508 protein	H92070	1,21
cyclin-dependent kinase 2	AB012305	1,21
ribosomal protein L5	BF214492	1,21
hypothetical protein 628	BC000974	1,21
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6 (17kD, B17)	NM_002493	1,21
dipeptidylpeptidase 8	NM_017743	1,21
hypothetical protein	NM_013313	1,21
cytochrome c oxidase subunit VI/c	NM_001867	1,21
hypothetical protein RG083M05.2	AI337901	1,21
a disintegrin and metalloproteinase domain 22	NM_021721	1,21
guanine nucleotide binding protein (G protein), gamma 7	NM_005145	1,21
diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)	NM_020548	1,21
tropomyosin 4	AI214061	1,21
Consensus includes gb:AK025152.1 /DEF=Homo sapiens cDNA: FLJ21499 fls, clone COL05634 /FEA=mRNA /DB_XREF=gi:10437612 /UG=Hs.306773 Homo sapiens cDNA: AK025152	NM_025152	1,21
KIAA0187 gene product	NM_014753	1,21
IKK-related kinase epsilon; inducible IkappaB kinase	AW340333	1,21
oviductal glycoprotein 1, 120kD (mucin 9, oviductin)	NM_002557	1,21
hypothetical protein FLJ12438	W60602	1,21
Ras-GTPase activating protein SH3 domain-binding protein 2	AU149503	1,21
polymerase (DNA directed), epsilon 3 (p17 subunit)	BC004170	1,21
Slah-interacting protein	AF275803	1,21
neuretic peptide receptor B/guanlylate cyclase B (atrial/natriuretic peptide receptor B)	AA565715	1,21
KIAA0337 gene product	NM_014786	1,21
hypothetical protein FLJ20080	NM_017657	1,21

cold inducible RNA binding protein
 super conserved receptor expressed in brain 3
 fibronectin 1
 LM protein (similar to rat protein kinase C-binding enigma)
 glycoprotein V (platelet)
 Consensus includes gb:AU134977 /FEA=EST /DB_XREF=gi:10995516 /DB_XREF=est:AU134977 /CLONE=PLACE1000926 /UG=Hs.322149 Human clone 137308 mRNA, par
 prefoldin 1
 rab6 GTPase activating protein (GAP and centrosome-associated)
 olfactory receptor, family 2, subfamily F, member 1
 phospholipidic acid phosphatase type 2B
 KIAA0438 gene product
 KIAA0792 gene product
 putative L-type neutral amino acid transporter
 extra spindle poles, S. cerevisiae, homolog of
 Xq28. 2000bp sequence contig. ORF
 hypothetical protein FLJ10597
 CDC28 protein kinase 1
 hypothetical protein FLJ20399
 COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 7B
 endothelin receptor type A
 hypothetical protein FLJ11289
 solute carrier family 28 (nucleoside transporters), member 2
 zinc finger protein-like 1
 DEAD/DEH (Asp-Glu-Ala-Asp/His) box polypeptide 34
 phospholipase C, epsilon 2
 nitrogen fixation cluster-like
 solute carrier family 1 (glutamate/neutral amino acid transporter), member 4
 FKBP58 binding protein 8 (35kD)
 DKE2P434F195 protein
 sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2
 enhancer of zeste homolog 1 (Drosophila)
 M-phase phosphoprotein 9
 protein tyrosine phosphatase, receptor type, F
 oxidative-stress responsive 1
 sudD (suppressor of bimD6, Aspergillus nidulans) homolog
 6.2 kd protein
 neuron-specific protein
 hypothetical protein FLJ20125
 Consensus includes gb:AW293356 /FEA=EST /DB_XREF=gi:6699992 /DB_XREF=est:LI-H-B12-eh1-c-11-Q-UI.s1 /CLONE=IMAGE:2727165 /UG=Hs.58220 Homo sapiens cDN
 selenophosphate synthetase 2
 inositol(myo)-1(or 4)-monophosphatase 1
 seven transmembrane domain orphan receptor
 gb:L14454.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398495 /FL=gb:L14454.1
 KIAA0467 protein
 tumor protein D52-like 2
 uncharacterized bone marrow protein BM039
 zinc finger protein 282
 small inducible cytokine subfamily B (Cys-X-Cys), member 6 (granulocyte chemolactic protein 2)
 Consensus includes gb:AL109705.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 73337. /FEA=mRNA /DB_XREF=gi:5689834 /UG=Hs.9997 Homo
 crooked neck protein (cm)
 Consensus includes gb:AA001552 /FEA=EST /DB_XREF=gi:1437017 /DB_XREF=est:ze46c08.s1 /CLONE=IMAGE:362030 /UG=Hs.59839 ESTs, Weakly similar to CGHUL7 c
 insulin induced protein 2
 hypothetical protein FLJ20259
 hypothetical protein FLJ10656
 huntingtin (Huntington disease)
 hypothetical protein FLJ12585
 granuln
 golgi SNAP receptor complex member 2
 hypothetical protein FLJ21865
 I-complex 10 (a murine tap homolog)
 mitogen-activated protein kinase kinase 2
 KIAA0650 protein
 male-specific lethal 3-like 1 (Drosophila)

protein phosphatase 2, regulatory subunit B (B56), gamma isoform	L42375	1.20
CGI-92 protein	NM_016035	1.20
polymerase (RNA) II (DNA directed) polypeptide A (220kD)	NM_000937	1.20
H2A histone family, member G	NM_021065	1.20
retinoblastoma-like 2 (p130)	BF110947	1.20
calcineurin binding protein 1	AB002328	1.20
HIV-1 Tat interactive protein 2, 30 kDa	BC002439	1.20
phosphoribosylformylglycinamide synthase (FGAR amidotransferase)	AL041328	1.20
UDP glycosyltransferase 1 family, polypeptide A8	NM_001072	1.20
persephin	NM_004158	1.20
hypothetical protein FLJ20298	NM_017750	1.20
linked to Surfeit genes in Fugu rubripes 2; LSFR2 gene 2	AI684664	1.20
solute carrier family 2 (facilitated glucose transporter), member 2	NM_000340	1.20
A kinase (PRKA) anchor protein 1	U34074	1.20
hypothetical protein FLJ17553	NM_024700	1.20
KIAA0842 protein	AB020849	1.20
zinc finger protein, Y-linked	NM_003411	1.20
KIAA1254 protein	AB033080	1.20
HpaII lny. fragments locus 9C	NM_022727	1.20
hypothetical protein FLJ20344	NM_017776	1.20
synaptosomal-associated protein, 23kD	U55936	1.20
inositol polyphosphate-5-phosphatase, 75kD	AK022846	1.20
aconitase 1, soluble	NM_002197	1.20
peroxisomal D3,D2-enoyl-CoA isomerase	NM_006117	1.20
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5 (13kD, B13)	AK022209	1.20
phospholipid scramblase 3	AB033924	1.20
nigamap homolog 1 (C. elegans)	AI783924	1.20
nucleic acid deoxyribose diphosphate linked moiety X-type motif 9	AW063371	1.20
KIAA0047 gene product	NM_024047	1.20
hypothetical protein FLJ20113	BE865244	1.20
E74-like factor 2 (ets domain transcription factor)	AL098714	1.20
suppressor of Ty 6 homolog (S. cerevisiae)	AF286223	1.20
mitogen-activated protein kinase 14	AI494587	1.20
KIAA0244 protein	AF100544	1.20
cathepsin D (lysosomal aspartyl protease)	BF430956	1.20
chromosome 18 open reading frame 1	NM_001909	1.20
synovial sarcoma, X breakpoint 1	NM_004338	1.20
hypothetical protein FLJ20477	BC001003	1.20
WD repeat domain 13	NM_017837	1.20
hypothetical protein DKFZp762P2111	AF158928	1.19
amiloride-sensitive calcium channel 3, testis	AU151157	1.19
DKFZP434D1335 protein	NM_020322	1.19
entyrin repeat domain 5	BC054966	1.19
HSPC052 protein	NM_022096	1.19
mucosa associated lymphoid tissue lymphoma translocation gene 1	NM_014150	1.19
glutathione reductase	AB026118	1.19
GMY2 ganglioside activator protein	NM_000637	1.19
KIAA0191 protein	X61094	1.19
ubiquitin 1	AI049862	1.19
Consensus includes gb:AW517716 /FEA=EST /DB_XREF=gi:7155798 /DB_XREF=est:xa86n03.x1 /CLONE=IMAGE:2776565 /UG=Hs.293472 ESTs	NM_016938	1.19
splicing factor (CC1.3)	AW517716	1.19
zinc finger protein 38, C3H type, homolog (mouse)	AI690947	1.19
KIAA0440 protein	NM_003407	1.19
glycoprotein A repetitions predominant	NM_015556	1.19
crystallin, alpha A	AP001748	1.19
protocadherin LKC	NM_005512	1.19
adenosine deaminase, RNA-specific	NM_017675	1.19
regulator of nonsense transcripts 1	NM_001111	1.19
sarcosomal vesicle protein 1	D66988	1.19
CGI-98 protein	NM_001612	1.19
pleomorphic adenoma gene-like 2	Z93241	1.19
hypothetical protein FLJ22418	NM_002857	1.19
v-yes-1 Yamaguchi sarcoma viral related oncogene homolog	NM_024626	1.19
	NM_002350	1.19

S100 calcium binding protein A13	NM_005878	1.19
peroxisomal fatty-acylated protein	NM_002857	1.19
hypothetical protein FLJ10120	NM_018001	1.19
hypothetical protein	NM_019049	1.19
pasillin	NM_002859	1.19
nuclear receptor coactivator 2	NM_006540	1.19
guanine nucleotide binding protein (G protein), alpha 2 polypeptide	NM_002073	1.19
ribosomal protein L26	AL121871	1.19
BTAF1 RNA polymerase II, B-TFIIID transcription factor-associated, 170 kD (Mat1 homolog, S. cerevisiae)	A001017	1.19
guanine nucleotide binding protein 10	NM_004125	1.19
Consensus includes gb-AA156865 /FEA=EST /DB_XREF=gi:1728480 /DB_XREF=est 20108.s1 /CLONE=IMAGE:502503 /UG=Hs.22370 Homo sapiens mRNA: cDNA DKFZp	AA156865	1.19
protein tyrosine phosphatase, receptor type, C-associated protein	NM_005608	1.19
FGF receptor activating protein 1	AL049281	1.19
Wiskott-Aldrich syndrome (eczema-thrombocytopenia)	U12707	1.19
lysine 3-monooxygenase/hypophan 5-monooxygenase activation protein, theta polypeptide	AA854017	1.19
protein kinase C, zeta	NM_002744	1.19
elastase 3B, pancreatic	NM_007352	1.19
urothrin (homologous to dystrophin)	NM_007124	1.19
serine/threonine kinase 25 (STE20 homolog, yeast)	NM_006374	1.19
synthrophin, beta 2 (dystrophin-associated protein A1, 59kD, basic component 2)	AA741303	1.19
chromobox homolog 3 (HP1 gamma homolog, Drosophila)	NM_016587	1.19
thyroid hormone receptor interactor 7	AF274949	1.19
chromobox homolog 6	NM_014292	1.19
I-kappa-B-interacting Ras-like protein 2	NM_017595	1.19
KIA0687 protein	A086285	1.19
bridging integrator 3	AK001289	1.19
hypothetical protein MGC3182	NM_024078	1.19
calcium channel, voltage-dependent, gamma subunit 2	AL022313	1.19
signal sequence receptor, alpha (transferrin-associated protein alpha)	AW006345	1.19
CD209 antigen	NM_021155	1.19
hypothetical protein PR00943	NM_018568	1.19
BCL2/adenovirus E1B 19kD interacting protein 3	NM_004052	1.19
muskelin 1, intracellular mediator containing kelch motifs	NM_013255	1.19
Q protein-binding protein CRFG	A081107	1.19
hypothetical protein FLJ13117	NM_023071	1.19
DKFZP586J1624 protein	NM_015537	1.19
hypothetical protein FLJ21212	NM_024642	1.19
cyclin M4	NM_020184	1.19
Consensus includes gb-AA885297 /FEA=EST /DB_XREF=gi:2994374 /DB_XREF=est 58103.s1 /CLONE=IMAGE:1461537 /UG=Hs.323567 CD36 antigen (collagen type I rec	AA885297	1.19
protein homologous to salivary proline-rich protein P-8	NM_012390	1.19
GDNF family receptor alpha 2	U87145	1.19
solute carrier family 24 (sodium/potassium/calcium exchanger), member 3	NM_020689	1.19
protein tyrosine phosphatase type IVA, member 1	BF576710	1.19
zinc finger protein 174	BC001161	1.19
hypothetical protein FLJ11128	NM_018332	1.19
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	M84630	1.19
KIA00923 protein	AU152563	1.19
cell matrix adhesion regulator	NM_005200	1.19
chromosome 8 open reading frame 2	T90773	1.19
Consensus includes gb-AK024450.1 /DEF=Homo sapiens mRNA for FLJ00042 protein, partial cds. /FEA=mRNA /GEN=FLJ00042 /PROD=FLJ00042 protein /DB_XREF=gi:104	AK024450	1.19
coplin III	AA541758	1.19
Consensus includes gb-AF228166.1 /DEF=Homo sapiens docking protein 1-like protein mRNA, partial cds. /FEA=mRNA /PROD=docking protein 1-like protein /DB_XREF=gi:71	AF229166	1.19
small nuclear ribonucleoprotein polypeptide C	NM_003093	1.19
PR domain containing 5	NM_018699	1.19
glycine C-acetyltransferase (2-smitho-3-ketobutyrate coenzyme A ligase)	NM_014281	1.19
nuclear autoantigenic sperm protein (histone-binding)	AY003362	1.19
Arg/Arg-interacting protein ArgBP2	NM_021069	1.19
muscleblind-like (Drosophila)	NM_021038	1.19
hypothetical protein FLJ13433	NM_022486	1.19
hypothetical protein FLJ10900	NM_018284	1.19
methyl-CpG binding domain protein 1	NM_015946	1.19
leukotriene A4 hydrolase	J02959	1.19
hypothetical protein FLJ20241	NM_011721	1.19

N-acetylserine amidohydrolase (acid ceramidase)-like
 dendritic cell protein
 RAD1 homolog (S. pombe)
 serine/threonine kinase 18
 tissue inhibitor of metalloproteinase 3 (Sorby fundus dystrophy, pseudoinflammatory)
 hypothetical protein FLJ20160
 HIV-1 Tat interactive protein, 60 kDa
 hypothetical protein FLJ10848
 fibronectin type 3 and SPRY domain-containing protein
 Fas (TNFRSF8) associated factor 1
 hypothetical protein dJ473B4
 hypothetical protein FLJ10706
 RacQ protein-like (DNA helicase O1-like)
 hypothetical protein MGC10772
 KIAA0874 protein
 cytochrome c oxidase subunit VIc pseudogene 1
 kraken-like
 Consensus includes gb:AL050035.1 /DEF=Homo sapiens mRNA; cDNA DKFp566H0124 (from clone DKFp566H0124) /FEA=mRNA /DB_XREF=gi:4884276 /UG=Hs.13313
 core promoter element binding protein
 cadherin 6, type 2, K-cadherin (fetal kidney)
 hypothetical protein FLJ10081
 TPA regulated locus
 somatostatin receptor 1
 katanin p80 (WD40-containing) subunit B 1
 follicle stimulating hormone receptor
 high-mobility group 20A
 I-box and leucine-rich repeat protein 4
 SRY (sex determining region Y)-box 1
 chemokine-like receptor 1
 Consensus includes gb:AL117381 /DEF=Human DNA sequence from clone RPS-357M17 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 3' p
 RNA cyclase homolog
 transient receptor potential channel 1
 KIAA0056 protein
 chromosome 21 open reading frame 45
 replication protein A3 (14kD)
 ATPase, H⁺ transporting, lysosomal (vacuolar proton pump), alpha polypeptide, 70kD, isoform 1
 KIAA0978 protein
 similar to hypothetical protein MNCb-2366
 tripartite motif-containing 31
 caltomer protein complex, subunit beta
 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2)
 cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)
 Notch homolog 3 (Drosophila)
 caspase 9, apoptosis-related cysteine protease
 guanine nucleotide binding protein (G protein), beta polypeptide 1
 Consensus includes gb:AU148611 /FEA=EST /DB_XREF=gi:11010132 /DB_XREF=est:AU148611 /CLONE=NT2RM400607 /UG=Hs.296832 Human pTR7 mRNA for repeltiv
 PTD002 protein
 gb:BC001788.1 /DEF=Homo sapiens. Similar to hypothetical protein from EUROMAGE 42353, clone MGC:2806, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypot
 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1)
 Deleted in split-hand/split-foot 1 region
 acetylserotonin O-methyltransferase-like
 KIAA0100 gene product
 Consensus includes gb:AV701318 /FEA=EST /DB_XREF=gi:10717648 /DB_XREF=est:AV701318 /CLONE=ADAAGD10 /UG=Hs.276268 ArgAbi-interacting protein ArgBP2
 CGL-87 protein
 Nijmegen breakage syndrome 1 (nibrin)
 JM1 protein
 hypothetical protein FLJ12785
 maternally expressed 3
 zinc finger protein 208
 gb:NM_000256.2 /DEF=Homo sapiens myosin-binding protein C, cardiac (MYBPC3), mRNA. /FEA=mRNA /GEN=MYBPC3 /PROD=protein C, cardiac /DB_XREF=gi:13325073
 menage a trois 1 (CAK assembly factor)
 aldo-keto reductase family 1, member C2 (dihydrodiol dehydrogenase 2; bile acid binding protein; 3-alpha hydroxysteroid dehydrogenase, type III)
 U05598

mevalonate kinase (mevalonic aciduria)
 hypothetical protein FLJ10604
 TNF receptor-associated factor 4
 vesicle-associated membrane protein 3 (collubrevin)
 protein tyrosine phosphatase, receptor type 1 (polypeptide (PTPR), interacting protein (liprin), alpha 1
 synaptobin, beta 1 (tyrosophilin-associated protein A1, 58kD, basic component 1)
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1
 HCDI protein
 trophobin
 chloride channel 3
 ubiquitin specific protease 11
 Consensus includes gb:AL35931.1 /FEA=EST /DB_XREF=gi:5768084 /DB_XREF=est:wr27b08.x1 /CLONE=IMAGE:248887 /UG=Hs.295923 seven in absentia (Drosophila) ho A1971256
 solute carrier family 20 (phosphate transporter), member 2
 F-box only protein 4
 Consensus includes gb:AL35931.1 /DEF=Homo sapiens mRNA full length insert cDNA done EUROMIMAGE 288936 /FEA=mRNA /DB_XREF=gi:8977886 /UG=Hs.48607 Homo sapiens
 high-mobility group (nonhistone chromosomal) protein 2
 KIAA1240 protein
 ribosomal protein L23a
 3-hydroxyisobutyryl-Coenzyme A hydrolase
 LIM domain kinase 1
 RNA binding protein S1, serine-rich domain
 hypothetical protein MGC3035
 nuclear transcription factor Y, gamma
 junctional adhesion molecule 1
 myosin, light polypeptide 4, alkali: atrial, embryonic
 cholinergic receptor, nicotinic, epsilon polypeptide
 hypothetical protein FLJ21047
 ADP-ribosylation factor GTPase activating protein 1
 Consensus includes gb:AW094068 /FEA=EST /DB_XREF=gi:6039220 /DB_XREF=est:xc26c06.x1 /CLONE=IMAGE:2585386 /UG=Hs.110630 Human BRCA2 region. mRNA
 hypothetical protein FLJ11026
 flightless 1 homolog (Drosophila)
 gene predicted from cDNA with a complete coding sequence
 mitochondrial translational initiation factor 2
 G protein-coupled receptor kinase-interactor 1
 melanin 1
 N-ethylmaleimide-sensitive factor
 hepatitis B virus x-interacting protein (9.6kD)
 cell division cycle 2-like 5 (cholinesterase-related cell division controller)
 cardiac-specific homeo box
 retinol binding protein 4, plasma
 gastric intrinsic factor (vitamin B synthesis)
 DiGeorge syndrome critical region gene DGS1: likely ortholog of mouse expressed sequence 2 embryonic lethal
 hypothetical protein MGC10882
 hypothetical protein FLJ21413
 TATA-binding protein-binding protein
 Consensus includes gb:AL052538 /FEA=EST /DB_XREF=gi:3308527 /DB_XREF=est:oz27g09.x1 /CLONE=IMAGE:1676608 /UG=Hs.31834 Homo sapiens done 25129 mRNA
 hypothetical protein FLJ13046 similar to exorphin 4
 peroxisomal biogenesis factor 14
 capping protein (actin filament), gelsolin-like
 cell division cycle 2-like 5 (cholinesterase-related cell division controller)
 KIAA0173 gene product
 Consensus includes gb:AL008730 /DEF=Human DNA sequence from PAC 487J7 on chromosome 6q21-22.1. Contains an unknown gene coding for three alternative mRNAs.
 protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calciueurin A alpha)
 oxidation resistance 1
 hypothetical protein FLJ20195
 KIAA0453 protein
 hypothetical protein FLJ13153
 Rho GTPase activating protein 8
 hypothetical protein FLJ13348
 gb:AF271775.1 /DEF=Homo sapiens DC48 mRNA, complete cds. /FEA=mRNA /PROD=DC48 /DB_XREF=gi:12006206 /UG=Hs.307093 Homo sapiens DC48 mRNA, complete
 tachykinin 3 (neuromedin K, neurokinin beta)
 timeless homolog (Drosophila)
 KIAA0682 gene product

S-adenosylhomocysteine hydrolase-like 1
 candidate phosphotransferase (nucleotide-nucleoside pyrophosphorylase (carboxylating))
 diphtheria toxin resistance protein required for diphtheria biosynthesis (Sacharomycetes)-like 1
 coagulation factor XI (plasma thromboplastin antecedent)
 HSPC128 protein
 chromosome 18 open reading frame 1
 early growth response 1
 hypothetical protein MGC2577
 hypothetical protein FLJ20203
 actin cross-linking factor
 bromodomain adjacent to zinc finger domain, 2B
 splicing factor, arginine/serine-rich 2
 major histocompatibility complex, class II, DP alpha 1
 hepatocyte growth factor (hepatopoietin A; scatter factor)
 hypothetical protein FLJ20511
 KIAA0233 gene product
 protymosin, alpha (gene sequence 2B)
 Consensus includes gb:AK022806.1 /DEF=Homo sapiens cDNA FLJ12744 fts, clone NT2RP2000715. /FEA=miRNA /DB_XREF=gi:10434418 /UG=Hs.139425 Homo sapiens cl
 Fc fragment of IgG, receptor, transporter, alpha
 KIAA1624 protein
 v-rei reticuloendotheliosis viral oncogene homolog A, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3, p65 (avian)
 BTB (POZ) domain containing 2
 Consensus includes gb:AK024680.1 /DEF=Homo sapiens cDNA: FLJ21207 fts, clone COL00382. /FEA=miRNA /DB_XREF=gi:10437268 /UG=Hs.306708 Homo sapiens cDNA:
 NME7
 hypothetical protein AF311304
 cytidine monophosphate-N-acetylneuraminic acid hydroxylase (CMP-N-acetylneuraminic acid monooxygenase)
 NAD kinase
 CAP-binding protein complex interacting protein 2
 hypothetical protein FLJ23469
 hypothetical protein FLJ21047
 Consensus includes gb:AW978196 /FEA=EST /DB_XREF=gi:8170484 /DB_XREF=est:EST391306 /UG=Hs.292713 ESTs, Moderately similar to ALLU1_HUMAN ALU SUBFAMI
 apoptosis inhibitor 5
 BCL2/adenovirus E1B 19kD interacting protein 2
 insulin receptor substrate 4
 insulin receptor family 5 (choline transporter), member 7
 solute carrier family 5 (choline transporter), member 7
 ubiquitin-protein isopeptide ligase (E3)
 A kinase (PRKA) anchor protein 8
 caspase recruitment domain protein 10
 frequently rearranged in advanced T-cell lymphomas 2
 keratin 10 (epidermolytic hyperkeratosis; keratosis palmaris et plantaris)
 hypothetical protein FLJ12881
 ELK3, ETS-domain protein (SRF accessory protein 2)
 phosphatidylinositol 4-kinase type II
 hypothetical protein FLJ10385
 sialoadhesin
 hypothetical protein FLJ11137
 zinc finger protein 42 (myeloid-specific retinoic acid responsive)
 KIAA0737 gene product
 forkhead box D3
 ribosomal protein L13a
 N-acylaminoacyl-peptide hydrolase
 hypothetical protein FLJ10883
 DiGeorge syndrome critical region gene 6 like
 suppressor of Ty 6 homolog (S. cerevisiae)
 Consensus includes gb:BF246115 /FEA=EST /DB_XREF=gi:11160133 /DB_XREF=est:60185408F1 /CLONE=IMAGE:4073921 /UG=Hs.8765 RNA helicase-related protein
 hypothetical protein FLJ10535
 cathelicidin antimicrobial peptide
 metastasis-associated 1-like 1
 Tcd37 homolog
 E74-like factor 3 (zeta domain transcription factor, epithelial-specific)
 insulin receptor substrate 2
 Consensus includes gb:AA071454 /FEA=EST /DB_XREF=gi:1578817 /DB_XREF=est:zm73c03.81 /CLONE=IMAGE:531268 /UG=Hs.258725 ESTs, Moderately similar to ATPC AA071454

FIGURE 4B.

gene	NCBI Accession Number	FOLD DECREASE
Consensus includes gb:AA292281 /FEA=EST /DB_XREF=gi:1940261 /DB_XREF=est:z51b03.s1 /CLONE=IMAGE:725837 /UG=Hs.181307 H3 histone, family 3A	AA292281	15.15
carleibulin	AA292281	13.38
ribosomal protein S11	AF1378708	10.86
ribosomal protein S19	BF680255	9.95
ribosomal protein, large P2	BC000023	8.60
Consensus includes gb:AW302047 /FEA=EST /DB_XREF=gi:5711724 /DB_XREF=est:x52108.x1 /CLONE=IMAGE:2763783 /UG=Hs.76230 ribosomal protein S10	AW302047	7.66
Consensus includes gb:L48784 /DEF=050 Homo sapiens cDNA /FEA=mRNA /DB_XREF=gi:1066715 /UG=Hs.182426 ribosomal protein S2	L48784	7.66
Consensus includes gb:AJ249377.1 /DEF=Homo sapiens partial mRNA for human Ig lambda light chain variable region, clone MB91 (331 bp). /FEA=mRNA (GENE=IGLV /P AJ249377	AJ249377	7.60
Consensus includes gb:D84143.1 /DEF=Human Immunoglobulin (mAb59) light chain V region mRNA, partial sequence. /FEA=mRNA (PROD=immunoglobulin light chain V D84143	D84143	6.93
ribosomal protein L27a	BE737027	6.41
emopamil binding protein (steroid isomerase)	N58463	5.40
carleibulin	AA910371	5.23
hypothetical protein PRO1843	NM_018507	4.91
Cluster Inci. A1201594:gc02h12.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-1708487 /clone_end=3 /gb=A1201594 /gi=3754200 /ug=Hs.239333 /len=591	A1201594	4.69
ribosomal protein S20	AF113008	4.57
C-terminal binding protein 1	BF984434	4.45
KIAA0908 protein	AA908765	4.42
ribosomal protein L27	BE312027	4.24
Consensus includes gb:AF044592 /DEF=Homo sapiens lymphocyte-predominant Hodgkins disease case 4 immunoglobulin heavy chain gene, variable region, partial cds // AF044592	AF044592	4.13
hypothetical protein FLJ21034	NM_024940	4.04
aminopeptidase puromycin sensitive	BG153399	4.00
immunoglobulin lambda locus	AF043566	3.96
Consensus includes gb:AJ239383.1 /DEF=Homo sapiens mRNA for immunoglobulin heavy chain variable region, ID 31. /FEA=mRNA (GENE=IGHV /PROD=immunoglobulin	AJ239383	3.81
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2	X93598	3.78
hypothetical protein FLJ12819	BE465032	3.67
lymphoid blast crisis oncogene	AF127481	3.63
Consensus includes gb:AV719355 /FEA=EST /DB_XREF=gi:10816507 /DB_XREF=est:AV719355 /CLONE=CLCEMB08 /UG=Hs.97109 ESTs	AV719355	3.59
alanine-glyoxylate aminotransferase 2-like 1	NM_031279	3.56
Cluster Inci. A1949010:wg36807.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2473332 /clone_end=3 /gb=A1949010 /gi=5741320 /ug=Hs.104036 /len=457	A1949010	3.49
PCTAIRE protein kinase 1	NM_006201	3.42
ADP-ribosylation factor 6	AA243143	3.42
lysine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide	AA502643	3.42
Consensus includes gb:BF973387 /FEA=EST /DB_XREF=gi:12340602 /DB_XREF=est:60224235F1 /CLONE=IMAGE:4330861 /UG=Hs.305989 Human DNA sequence	BF973387	3.37
Q protein-coupled receptor kinase 6	BG423052	3.33
M10098 Human 18S rRNA sequence, length 1968 bases, middle target bases 647-1292	M10098	3.24
GM2 ganglioside activator protein	X61094	3.24
dystrophia myotonic-containing WD repeat motif	L19267	3.24
collagen, type I, alpha 1	AJ743521	3.22
cathepsin S	BC026842	3.19
transition initiation factor IF2	AB016284	3.16
hypothetical protein FLJ22965	NM_022101	3.14
coactivator-associated arginine methyltransferase-1	AF116815	3.13
JTV1 gene	NM_005733	3.10
RA86 Interacting, kinesin-like (raklasein 6)	NM_018333	3.06
hypothetical protein FLJ20666	NM_005696	3.04
isodrate dehydrogenase 1 (NADP+), soluble	NM_018333	3.03
Consensus includes gb:A1524687 /FEA=EST /DB_XREF=gi:4438822 /DB_XREF=est:h12a07.x1 /CLONE=IMAGE:2118036 /UG=Hs.57969 phenylalanine-tRNA synthetase A1524687	A1524687	3.02
Consensus includes gb:AL138178 /DEF=Human DNA sequence from clone RP3-322L4 on chromosome 6. Contains the SOX4 gene for SRY (sex determining region Y)-box AL138178	AL138178	3.02
nucleolar protein 4	NM_003787	2.98
ribosomal protein L38	AW303136	2.91
ribosomal protein L38	BC000603	2.90
Consensus includes gb:AW090043 /FEA=EST /DB_XREF=gi:6047387 /DB_XREF=est:td01c05.x1 /CLONE=IMAGE:2592488 /UG=Hs.326464 Homo sapiens cDNA: FLJ21 AW090043	AW090043	2.88
Consensus includes gb:AW971415 /FEA=EST /DB_XREF=gi:8161260 /DB_XREF=est:EST3383504 /UG=Hs.165337 ESTs	AW971415	2.87
J04423 E coli bcd gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423	2.85
Consensus includes gb:AA890010 /FEA=EST /DB_XREF=gi:3016889 /DB_XREF=est:aj89h08.s1 /CLONE=IMAGE:1403679 /UG=Hs.50785 SEC22, vesicle trafficking prot AA890010	AA890010	2.83
gb:Z25437.1 /DEF=H sapiens protein-tyrosine kinase gene, complete CDS. /FEA=mRNA (PROD=protein-tyrosine kinase /DB_XREF=gi:405752 /FL=gb.Z25437.1	Z25437	2.81
prefoldin 4	NM_002823	2.81
KIAA1046 protein	NM_014928	2.81
ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1A (110/116kD)	NM_005177	2.81

134

hypothetical protein FLJ10159 NM_018013 2.80
Consensus includes gb:BG109746 /FEA=EST /DB_XREF=gi:12603252 /DB_XREF=est:602280883F1 /CLONE=IMAGE:4368381 /UG=Hs.325625 Homo sapiens clone 239 BG109746 2.79
HIV-1 rev binding protein 2 A912583 2.76
acidic epididymal glycoprotein-like 1 X95238 2.75
hypothetical protein FLJ12619 AL136632 2.74
hypothetical protein FLJ14107 NM_025026 2.73
Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2393-2682 of J04423 /LEN=1114 (-5'..-M..-3' represent transcript region) J04423 2.73
E74-like factor 4 (ets domain transcription factor) NM_001421 2.71
ribonuclease P, 40KD subunit NM_006638 2.71
transducin (beta)-like 1 AAT24134 2.70
cytochrome c oxidase subunit Vb A557312 2.70
RAB5B, member RAS oncogene family AF267863 2.68
RAP2B, member of RAS oncogene family NM_002886 2.67
ATP-binding cassette, sub-family D (ALD), member 3 NM_002858 2.67
Consensus includes gb:AK023724.1 /DEF=Homo sapiens cDNA, FLJ22071 fls, clone HEP11691, /FEA=mrna /DB_XREF=gi:10438333 /UG=Hs.326248 Homo sapiens cDNA AK023724 2.67
soluble carrier family 21 (organic anion transporter), member 6 AB026257 2.65
Consensus includes gb:AW971134 /FEA=EST /DB_XREF=gi:8160979 /DB_XREF=est:EST383221 /UG=Hs.292245 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAM1 AW971134 2.64
orphan seven-transmembrane receptor, chemokine related NM_016557 2.61
Consensus includes gb:AL050065.1 /DEF=Homo sapiens mRNA, cDNA DKFZp566M043 (from clone DKFZp566M043), /FEA=mrna /DB_XREF=gi:4884295 /UG=Hs.2125 AL050065 2.61
PAX transcription activation domain interacting protein 1 like A357401 2.60
cytoskeleton-associated protein 4 NM_006825 2.59
KIAA0653 protein, B7-like protein AF289028 2.59
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 AF200738 2.59
epithelial membrane protein 3 NM_001425 2.59
ribosomal protein L37a BE85772 2.59
Consensus includes gb:A345238 /FEA=EST /DB_XREF=gi:4092444 /DB_XREF=est:db1807.x1 /CLONE=IMAGE:2060725 /UG=Hs.111334 ferritin, light polypeptide A345238 2.58
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9KD A252582 2.57
Bloom syndrome NM_000057 2.57
cytochrome P450 isoform 4F12 NM_023944 2.56
H3 histone family, member C NM_003531 2.56
hypothetical protein FLJ22009 NM_024745 2.56
glycophorin E NM_002102 2.55
hypothetical protein FLJ10298 NM_018050 2.55
endomucin-1 NM_016241 2.55
cyclin G2 AW134535 2.55
hexokinase 2 A1761561 2.54
polymerase (DNA directed), eta NM_006502 2.53
culin 5 NM_003478 2.53
PI-3-kinase-related kinase SMG-1 BE000837 2.52
helicase KIAA0054 NM_014104 2.50
M10098 Human 18S rRNA gene, complete (-5'..M..3' represent transcript regions 5 prime, Middle, and 3 prime respectively) NM_014877 2.50
KIAA0869 protein M10098 2.50
hypothetical protein FLJ20897 NM_015377 2.49
gamma-aminobutyric acid (GABA) A receptor, alpha 5 A335509 2.49
translation initiation factor IF2 BF966183 2.49
SCAM domain-containing 2 BE138647 2.48
inhibin, beta C AF244812 2.47
M10098 Human 18S rRNA sequence, length 1969 bases, 3 prime target bases 1293-1938 NM_005538 2.47
cysteine-rich motor neuron 1 M10098 2.47
mitochondrial ribosomal protein S12 BG546884 2.47
hypothetical protein FLJ10357 R88573 2.47
mamian-binding lectin serine protease 1 (C41C2 activating component of Re-active factor) NM_018071 2.47
thiopurine S-methyltransferase U12387 2.45
sorting nexin 4 AA523345 2.45
golgi associated, gamma adaptin ear containing, ARF binding protein 1 AW001443 2.44
Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2772-3004 of J04423 /LEN=1114 (-5'..-M..-3' represent transcript region) J04423 2.44
methionine adenosyltransferase II, alpha AW301861 2.43
Consensus includes gb:AK022473.1 /DEF=Homo sapiens cDNA FLJ12411 fls, clone MAMMA1002964, /FEA=mrna /DB_XREF=gi:10433882 /UG=Hs.256722 Homo sapiens cDNA AK022473 2.43
eukaryotic translation initiation factor 5A A3383940 2.42
tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein NM_021980 2.42
J04423 E coli bioB gene biotin synthetase (-5'..-M..-3' represent transcript regions 5 prime, Middle, and 3 prime respectively) J04423 2.42
SH3-domain GRB2-like 3 AF036269 2.42

hypothetical protein FLJ13078
 serine/threonine-protein kinase PRP4 homolog
 herak1, BCL2 interacting protein (contains only BH3 domain)
 Consensus includes gb:AK021505.1 /DEF=Homo sapiens cDNA FLJ11443 fls, clone HEMBA1001330 /FEA=mRNA /DB_XREF=gi:10432701 /UG=Hs.297945 Homo sapie
 genome pathogenesis-related protein
 artemis protein
 phosphodiesterase 10A
 ubiquitin specific protease 15
 TGF-beta-induced factor (TAFI family homeobox)
 PRO0478 protein
 artemis protein
 Consensus includes gb:A015947 /FEA=EST /DB_XREF=gi:5635802 /DB_XREF=ast:wg9601.x1 /CLONE=IMAGE:2378096 /UG=Hs.28212 ESTs
 triple functional domain (PTPRF interacting)
 thromboxane A2 receptor
 KIAA1655 protein
 Consensus includes gb:BF942161 /FEA=EST /DB_XREF=gi:12359481 /DB_XREF=ast:nae87g10.x1 /CLONE=IMAGE:4118994 /UG=Hs.302797 ESTs
 DnaJ (Hsp40) homolog, subfamily A, member 1
 hypothetical protein FLJ22479
 transforming, acidic coiled-coil containing protein 2
 KIAA0446 gene product
 DKFZP947E1010 protein
 Consensus includes gb:AL157484.1 /DEF=Homo sapiens mRNA, cDNA DKFZP762M127 (from clone DKFZP762M127) /FEA=mRNA /DB_XREF=gi:7018527 /UG=Hs.224E
 kinesin 1 (kinesin receptor)
 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
 DEAD/DH (Asp-Glu-Ala-Asp/His) box binding protein 1
 hypothetical protein FLJ23548
 MAD, mothers against decapentaplegic homolog (Drosophila) interacting protein, receptor activation anchor
 Consensus includes gb:BC004344.1 /DEF=Homo sapiens, clone IMAGE:3633354, mRNA, partial cds /FEA=mRNA /PROD=Unknown (protein for IMAGE:3633354) /DB_XI
 Consensus includes gb:R33964 /FEA=EST /DB_XREF=gi:789822 /DB_XREF=ast:y74c03.1 /CLONE=IMAGE:135460 /UG=Hs.286681 Homo sapiens cDNA FLJ11022 fls
 prostate derived STE20-like kinase PSK
 putative protein O-mannosyltransferase
 Consensus includes gb:AK021440.1 /DEF=Homo sapiens cDNA FLJ11378 fls, clone HEMBA1000456 /FEA=mRNA /DB_XREF=gi:10432825 /UG=Hs.6937
 Consensus includes gb:AL080180.1 /DEF=Homo sapiens mRNA, cDNA DKFZP434M054 (from clone DKFZP434M054) /FEA=mRNA /DB_XREF=gi:5282622 /UG=Hs.274E
 uncharacterized hypothetical protein HT011
 S-adenosylmethionine decarboxylase 1
 Consensus includes gb:BE786164 /FEA=EST /DB_XREF=gi:10207362 /DB_XREF=ast:60147273F1 /CLONE=IMAGE:3877146 /UG=Hs.80285 Homo sapiens mRNA, cD
 Consensus includes gb:BC005385.1 /DEF=Homo sapiens, clone IMAGE:3829438, mRNA, partial cds /FEA=mRNA /PROD=Unknown (protein for IMAGE:3829438) /DB_XI
 Consensus includes gb:AW971254 /FEA=EST /DB_XREF=gi:8161059 /DB_XREF=ast:EST383343 /UG=Hs.178433 ESTs
 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A
 Consensus includes gb:AF184963.1 /DEF=Homo sapiens tumor antigen NA88-A pseudogene, complete sequence /FEA=mRNA /DB_XREF=gi:5901726 /UG=Hs.306576 f
 immunoglobulin heavy constant mu
 actin-related protein 3-beta
 CDC14 cell division cycle 14 homolog A (S. cerevisiae)
 DEAD/DH (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72KD)
 PABP-interacting protein 2
 DnaJ (Hsp40) homolog, subfamily B, member 9
 alcohol dehydrogenase 1C (class I), gamma polypeptide
 ectopic viral integration site 2A
 Consensus includes gb:A126492 /FEA=EST /DB_XREF=gi:3595006 /DB_XREF=ast:qd82n06.x1 /CLONE=IMAGE:1736027 /UG=Hs.104258 Homo sapiens mRNA, exon 1
 N-myristoyltransferase 2
 J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
 phosphatidylcholine
 hypothetical protein FLJ22558
 peptidyl-prolyl isomerase G (cyclophilin G)
 KIAA0469 gene product
 putative lymphocyte G0/G1 switch gene
 Consensus includes gb:BE930512 /FEA=EST /DB_XREF=gi:10456588 /DB_XREF=ast:RC6-GN0071-160800-021-D01 /UG=Hs.168732 ESTs
 A kinase (PRKA) anchor protein 1
 aldehyde dehydrogenase 4 family, member A1
 Consensus includes gb:BF573849 /FEA=EST /DB_XREF=gi:1647561 /DB_XREF=ast:602132053F1 /CLONE=IMAGE:4271340 /UG=Hs.98343 ESTs, Weakly similar to A
 ATPase, Class I, type BB, member 1
 Consensus includes gb:A101158 /FEA=EST /DB_XREF=ast:we10109.x1 /CLONE=IMAGE:2340713 /UG=Hs.6550 Homo sapiens cDNA: FLJ23227 AT701156
 stress 70 protein chaperone, microsome-associated, 60kD
 AIT701156

cytochrome P450, subfamily I1C (mephenytoin 4-hydroxylase), polypeptide 19	X65962	2.08
choline/ethanolaminephosphotransferase	NM_006090	2.07
KIAA0304 gene product	AF105279	2.07
endase 1, (alpha)	U88968	2.07
H2A histone family, member X	H51429	2.07
ALEX3 protein	NM_016607	2.07
Consensus includes gb:AK026825.1 /DEF=Homo sapiens cDNA: FLJ23172 fis, clone LNG10005. /FEA=mRNA /DB_XREF=gi:10439771 /UG=Hs.306885 Homo sapiens cDNA: AK026825	NM_017644	2.07
hypothetical protein FLJ20059	NM_004982	2.07
potassium inwardly-rectifying channel, subfamily J, member 8	L28650	2.07
early lymphoid activation protein	X60821	2.06
KIAA0874 protein	NM_001052	2.06
somatostatin receptor 4	NM_025235	2.06
antikyrase, TRF1-interacting ankryrin-related ADP-ribose polymerase 2	BF002659	2.06
immunoglobulin heavy constant gamma 3 (G3m marker)	NM_012452	2.05
transmembrane activator and CAML interactor	NM_012415	2.05
RAD54, S. cerevisiae, homolog of B	AW770998	2.05
polymerase (RNA) II (DNA directed) polypeptide B (140kD)	N85418	2.05
FK506 binding protein 8 (38kD)	BG281322	2.05
translation initiation factor IF2	AL050154	2.05
Consensus includes gb:AL050154.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586L0120 (from clone DKFZp586L0120) /FEA=mRNA /DB_XREF=gi:4884366 /UG=Hs.985 AL050154	AL147017	2.05
Consensus includes gb:AU147017 /FEA=EST /DB_XREF=gi:11008538 /DB_XREF=est:AU147017 /CLONE=HEMBB1002152 /UG=Hs.301905 Homo sapiens cDNA: FLJ141	AL046992	2.05
G protein-coupled receptor 1	NM_006799	2.05
protease, serine, 21 (testisin)	NM_018114	2.05
hypothetical protein FLJ10496	NM_002785	2.05
pregnancy specific beta-1-glycoprotein 11	AI221318	2.04
RAN binding protein 1	AK022213	2.04
hypothetical protein FLJ12151	AK022174	2.04
Consensus includes gb:AK022174.1 /DEF=Homo sapiens cDNA: FLJ12112 fis, clone MAMMA1000043. /FEA=mRNA /DB_XREF=gi:10433510 /UG=Hs.288793 Homo sapi	AK026674	2.04
transcription factor 4	AI989477	2.04
SRY (sex determining region Y)-box 4	AC004460	2.03
Consensus includes gb:AC004460 /DEF=Homo sapiens PAC clone RPS-1086D14 /FEA=CDS /DB_XREF=gi:2981263 /UG=Hs.307352 Homo sapiens PAC clone RPS-1086	AK023918	2.03
Consensus includes gb:AK023918.1 /DEF=Homo sapiens cDNA: FLJ13356 fis, clone THYRO1000988. /FEA=mRNA /DB_XREF=gi:10436003 /UG=Hs.288489 Homo sapi	NM_014822	2.03
SEC24 related gene family, member D (S. cerevisiae)	NM_018393	2.03
hypothetical protein FLJ11338	AA860806	2.03
checkpoint suppressor 1	AF070647	2.03
Consensus includes gb:AF070647.1 /DEF=Homo sapiens clone 24438 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283921 /UG=Hs.124126 Homo sapiens clone 24438	AI659160	2.02
translin	AA295257	2.02
neurophilin 2	NM_018441	2.02
crystallin-rich motor neuron 1	NM_005620	2.02
S100 calcium binding protein A11 (calgizzarin)	NM_017648	2.02
gb:NM_017648.1 /DEF=Homo sapiens hypothetical protein FLJ20063 (FLJ20063), mRNA. /FEA=mRNA /GEN=FLJ20063 /PROD=hypothetical protein FLJ20063 /DB_XREF=	AF068482	2.02
PC4 and SFRS1 interacting protein 1	S74639	2.02
immunoglobulin heavy constant mu	AW301808	2.02
Consensus includes gb:AW301808 /FEA=EST /DB_XREF=gi:5711483 /DB_XREF=est:rs5611.1 /CLONE=IMAGE:2764172 /UG=Hs.150551 ESTs, Weakly similar to ALU	BF588511	2.01
ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome)	BG338608	2.01
aductin 2 (beta)	NM_018810	2.01
thyroid hormone receptor-associated protein, 95-kD subunit	AI206560	2.01
hypothetical protein PRO1942	AW772123	2.01
HCF-binding transcription factor Zhangfei	AK023368	2.00
protein phosphatase 2, regulatory subunit B (B56), gamma isoform	AK001699	2.00
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	AF080895	2.00
F-box only protein 21	NM_022335	2.00
gb:AF080895.1 /DEF=Homo sapiens clone HQ0117 PRO0117 mRNA, complete cds. /FEA=mRNA /PROD=PRO0117 /DB_XREF=gi:6690166 /UG=Hs.283919 Homo sapi	NM_003755	2.00
hypothetical protein PRO2849	L14561	2.00
Epstein-Barr virus induced gene 3	NM_015271	2.00
ATPase, Ca++ transporting, plasma membrane 1	AB011122	2.00
tripartite motif-containing 2	X78262	2.00
brain-specific angiogenesis inhibitor 3	NM_013984	2.00
Consensus includes gb:X78262.1 /DEF=H sapiens mRNA for TRES. /FEA=mRNA /DB_XREF=gi:587440 /UG=Hs.302178 H.sapiens mRNA for TRES	NM_005377	1.99
neuregulin 2	AB038463	1.99
v-myc myelocytomatosis viral oncogene homolog 2 (avian)	NM_012482	1.99
calpain 9 (nCL-4)		
zinc finger protein 281		

hypothetical protein LOC57187	BG403871	1.99
hypothetical protein FLJ13166	NM_025003	1.99
gastric inhibitory polypeptide receptor	NM_000164	1.99
activation-induced cytidine deaminase	NM_020861	1.99
caspase 8, apoptosis-related cysteine protease	BF439983	1.99
hypothetical protein FLJ20837	NM_017864	1.99
dictyof homolog 2 (Xenopus laevis)	NM_014421	1.99
lipopolysaccharide specific response-68 protein	NM_018678	1.99
protein phosphatase, EF hand calcium-binding domain 2	NM_006239	1.98
hypothetical protein FLJ14348	NM_025028	1.98
membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for; beta polypeptide)	NM_000139	1.98
Escherichia coli (REF=J04423)/DEF=E coli bdc protein corresponding to nucleotides 4609-4883 of J04423/LEN=777 (-5 and -3 represent transcript regions 5 prime and 3 J04423)	J04423	1.98
Consensus includes gb:BF035279 /FEA=EST /DB_XREF=gi:10743008 /DB_XREF=est:601457185F1 /CLONE=IMAGE:3860633 /UG=Hs.20325 ESTs. Moderately similar to AK027231	AK027231	1.98
KIAA1102 protein	NM_002408	1.98
mannosyl (alpha-1,6)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	AA047234	1.98
onitidine decarboxylase anilzyme inhibitor	NM_015896	1.98
BLU protein	NM_005028	1.98
phosphatidylinositol-4-phosphate 5-kinase, type II, alpha	NM_016415	1.98
clone FLB316	BC026566	1.97
thyroid hormone receptor-interactor 11	Y13710	1.97
small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated	NM_025002	1.97
hypothetical protein FLJ13162	BE550452	1.97
Homer, neuronal immediate early gene, 1B	NM_025028	1.97
hypothetical protein FLJ14310	NM_043804	1.97
Consensus includes gb:U43804.1 /DEF=Human unidentified mRNA, partial sequence. /FEA=mRNA /DB_XREF=gi:1171236 /UG=Hs.159801 Human unidentified mRNA, pt U43804	NM_000198	1.97
hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 2	U55968	1.97
lipin 2	AL050122	1.97
Consensus includes gb:AL050122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586E121 (from clone DKFZp586E121) /FEA=mRNA /DB_XREF=gi:4884330 /UG=Hs.2745	AL050122	1.97
gb:U96291.1 /DEF=Homo sapiens Ig kappa light chain variable region (VKII-A23) mRNA, complete cds. /FEA=mRNA /GEN=VKII-A23 /PRODB=ig kappa light chain variable region U96291	NM_022164	1.97
PAGECSL	AL044175	1.97
patched homolog (Drosophila)	A1734228	1.97
WW domain binding protein 4 (fornin binding protein 21)	NM_004540	1.97
neural cell adhesion molecule 2	NM_018041	1.97
hypothetical protein FLJ10254	AA910614	1.96
ubiquitin-conjugating enzyme E2i (UBCS homolog, yeast)	NM_021093	1.96
peptide YY, 2 (seminalplasmin)	AF116696	1.96
DnaJ (Hsp40) homolog, subfamily C, member 8	AK025933	1.96
KIAA0692 protein	A1769637	1.96
UDP-glucuronic acid/UDP-N-acetylglucosamine dual transporter	NM_017659	1.96
hypothetical protein FLJ20084	BF668960	1.96
beta-amyloid binding protein precursor	AL133846	1.96
glucocorticoid modulatory element binding protein 2	NM_000130	1.96
coagulation factor V (proaccelerin, labile factor)	N52532	1.96
KIAA0256 gene product	AK000864	1.96
Consensus includes gb:AK000864.1 /DEF=Homo sapiens cDNA FLJ10002 /is. clone HEMBA1000046. /FEA=mRNA /DB_XREF=gi:7021188 /UG=Hs.296522 Homo sapiens	NM_022915	1.96
mitochondrial ribosomal protein L44	NM_022219	1.95
Consensus includes gb:AK022219.1 /DEF=Homo sapiens cDNA FLJ12157 /is. clone MAMMA1000500. /FEA=mRNA /DB_XREF=gi:10433569 /UG=Hs.98812 Homo sapiens	BG029530	1.95
junonil homolog (mouse)	NM_018953	1.95
homeo box C5	U52896	1.95
cAMP responsive element binding protein-like 1	NM_014274	1.95
Alu-binding protein with zinc finger domain	NM_014809	1.95
KIAA0318 gene product	NM_002078	1.95
golgi autoantigen, golgin subfamily a, 4	BC001265	1.95
hypothetical protein dJ482023.2	NM_004696	1.94
soluble carrier family 16 (monocarboxylic acid transporters), member 4	AF220028	1.94
tripartite multi-containing 5	NM_021999	1.94
integral membrane protein 2B	NM_000052	1.94
ATPase, Cu++ transporting, alpha polypeptide (Menkes syndrome)	NM_004702	1.94
cyclin E2	X75208	1.94
EphB3	NM_017667	1.94
hypothetical protein FLJ20097	NM_012328	1.94
DnaJ (Hsp40) homolog, subfamily B, member 9	NM_006467	1.94
polymerase (RNA) III (DNA directed) (32XO)		

aldehyde dehydrogenase 1 family, member B1
 Consensus includes gb:AA017721 /FEA=EST /DB_XREF=gi:1479910 /DB_XREF=est:z639f11.31 /CLONE=IMAGE:361389 /UG=Hs.49.117 Homo sapiens mRNA; cDNA
 nuclear transcription factor Y, alpha
 calcitonin receptor
 tubby like protein 3
 hypothetical protein FLJ11328
 golgi associated, gamma adaptin ear containing, ARF binding protein 2
 Consensus includes gb:AK023783.1 /DEF=Homo sapiens cDNA FLJ13721 fls, done PLACE2000450. /FEA=mRNA /DB_XREF=gi:10435820 /UG=Hs.289035 Homo sapiens
 succinate dehydrogenase complex, subunit C, intragolgi membrane protein, 15kD
 hypothetical protein FLJ20019
 KIAA1110 protein
 Consensus includes gb:BC049571 /FEA=EST /DB_XREF=gi:13457287 /DB_XREF=est:802540396F1 /CLONE=IMAGE:4671599 /UG=Hs.289068 Homo sapiens cDNA FLJ
 zinc finger protein 43 (HIF6)
 Fc fragment of IgG, low affinity IIb, receptor for (CD16)
 hypothetical protein similar to small G proteins, especially RAP-2A
 synovial sarcoma translocation gene on chromosome 18-like 1
 Consensus includes gb:AW576457 /FEA=EST /DB_XREF=est:UH-HF-BR0p-slj-a-01-Q-U1.s1 /CLONE=IMAGE:3074658 /UG=Hs.20952 Homo sapiens
 prostate differentiation factor
 KIAA1827 protein
 trichothiodystrophy syndrome 1
 dynamin 2
 dopamine receptor D4
 Consensus includes gb:AK025095.1 /DEF=Homo sapiens cDNA: FLJ21442 fls, done COL04429, highly similar to HSA237839 Homo sapiens mRNA for hypothetical protein
 NG37 protein
 protein kinase, cAMP-dependent, regulatory, type II, alpha
 hypothetical protein DKFZp34D0421
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 2
 phospholamban
 Consensus includes gb:AV684859 /FEA=EST /DB_XREF=gi:10286722 /DB_XREF=est:AV684859 /CLONE=GKCDJAO3 /UG=Hs.251334 ESTs, Moderately similar to ALU6
 glutamate receptor, ionotropic, kainate 4
 male germ cell-associated kinase
 fatty-acyl-Coenzyme A ligase, long-chain 3
 Fc fragment of IgE, high affinity I, receptor for: alpha polypeptide
 neuronal Shc
 hypothetical protein FLJ23412
 transmembrane protein 5
 sorting nexin 15
 G-2 and S-phase expressed 1
 Consensus includes gb:AW970584 /FEA=EST /DB_XREF=gi:8160429 /DB_XREF=est:EST382685 /UG=Hs.291033 ESTs
 KIAA0480 gene product
 Consensus includes gb:AW451711 /FEA=EST /DB_XREF=gi:6992487 /DB_XREF=est:UH-B13-ale-c-02-Q-U1.s1 /CLONE=IMAGE:2736386 /UG=Hs.313760 ESTs
 protein tyrosine kinase 9
 monoamine oxidase A
 homeo box A10
 nuclear factor I/B
 KIAA0415 gene product
 GABA(A) receptor-associated protein like 1
 hypothetical protein PRQ0082
 bone morphogenetic protein receptor, type IB
 Consensus includes gb:A1732770 /FEA=EST /DB_XREF=gi:5053383 /DB_XREF=est:z78d05.x5 /CLONE=IMAGE:809865 /UG=Hs.328688 ESTs, Moderately similar to AL
 pleomorphic adenoma gene 1
 KIAA0341 protein
 Consensus includes gb:AL312703.1 /DEF=Homo sapiens mRNA; cDNA DKFZp667H046 (from clone DKFZp667H046) /FEA=mRNA /GEN=DKFZp667H046 /PROD=hypoth
 Consensus includes gb:A1734156 /FEA=EST /DB_XREF=gi:5055269 /DB_XREF=est:mc79004.y5 /CLONE=IMAGE:783559 /UG=Hs.172895 ESTs
 tripeptidyl peptidase II
 glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV)
 intracellular hyaluronan-binding protein
 LIM domain protein
 mitogen-activated protein kinase 8
 SRY (sex determining region Y)-box 21
 hypothetical protein SP192
 Wilms tumor 1

laminin, beta 1	NM_002291.1
Consensus includes gb:A002715 /FEA=EST /DB_XREF=gj:3200129 /DB_XREF=est:an20612.s1 /CLONE=IMAGE:1699223 /UG=Hs.130315 ESTs	
general transcription factor IIH, polypeptide 3 (34kD subunit)	A002715
pallid zinc finger protein from EUROMAGE 566589	A0589458
killer cell lectin-like receptor subfamily D, member 1	NM_021220
doublesex and mab-3 related transcription factor 1	NM_007334
estrogen receptor 2 (ER beta)	NM_021951
Consensus includes gb:AV301806 /FEA=EST /DB_XREF=gj:6711483 /DB_XREF=est:c56a.11.x1 /CLONE=IMAGE:2764172 /UG=Hs.150551 ESTs, Weakly similar to ALU AW301806	AF060555
similar to rat nuclear ubiquitous casein kinase 2	AW515443
antigen identified by monoclonal antibody MRC OX-2	H23979
glycoprotein IX (platelet)	NM_000174
B-cell CLL/lymphoma 7A	A0950380
basic helix-loop-helix domain containing, class B, 3	BE857425
IIMP (inosine monophosphate) dehydrogenase 1	NM_000983
PC3-9g protein	NM_022488
cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)	AF115544
matrix metalloproteinase 25	NM_022716
death-associated protein kinase 2	AK026801
Consensus includes gb:AF222691.1 /DEF=Homo sapiens Alu repeat (LNX1) mRNA sequence. /FEA=mRNA /DB_XREF=gj:12655850 /UG=Hs.307008 Homo sapiens Alu AF222691	AF222691
f-box and leucine-rich repeat protein 11	AK024505
RAB3A, member RAS oncogene family	AA98824.1
CED-4 protein	AK023968
Consensus Includes gb:A0021609.1 /DEF=Homo sapiens cDNA FLJ11547 fls, clone HEMBA1002934, /FEA=mRNA /DB_XREF=gj:1043282 /UG=Hs.306605 Homo sapie	AK021609
ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome	NM_007125
nephrosis 1, congenital, Finnish type (nephrt)	NM_004846
HSPC039 protein	AF119875
BRCA1 associated protein	AL042733
3-phosphoinositide dependent protein kinase-1	AK026947
Consensus includes gb:BE872313 /FEA=EST /DB_XREF=gj:10032854 /DB_XREF=est:7a59b10.x1 /CLONE=IMAGE:3223003 /UG=Hs.34054 Homo sapiens cDNA: FLJ22	BE872313
Consensus includes gb:AL138105 /DEF=Human DNA sequence from clone RP4-670F13 on chromosome 1q42.2-4.3. Contains an endonase 1, (alpha) (ENO1) pseudogene,	AL138105
PQZ domain containing guanine nucleotide exchange factor(GEP1)	AL17397
hypothetical protein FLJ23142	NM_024593
a disintegrin-like and metalloprotease (repotysin type) with thrombospondin type 1 motif, 1	AK023795
KIAA1045 protein	AB028968
craniofacial development protein 1	D85939
G protein-coupled receptor 37 (endothelin receptor type B-like)	T16257
CS box-containing WD protein	BG250658
KIAA1885 protein	NM_018263
hypothetical protein FLJ22748	NM_024785
J04423 E coli bioC protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423
signal transduction protein (SH3 containing)	AB001467
KIAA0358 protein	A1272899
hypothetical protein FLJ13448	NM_025147
melanocortin 1 receptor (alpha melanocyte stimulating hormone receptor)	BG034872
Consensus includes gb:U63542.1 /DEF=Human putative FAP protein mRNA, partial cds. /FEA=mRNA /PROD=FAP protein /DB_XREF=gj:1890846 /UG=Hs.166101 Humar	U63542
likely ortholog of mouse polydrom	NM_024500
Heterogeneous nuclear ribonucleoprotein H3 (2H9)	NM_021644
kangal, 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (P2 leukocyte antigen, antigen detected by monoclonal and antibody lA4))	NM_002231
carbonic phosphatase B1 (tissue)	NM_001871
solute carrier family 13 (sodium/sulfate symporters), member 1	AF260824
hypothetical protein FLJ12873	NM_025133
J04423 E coli bioD gene dehalobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423
KIAA0584 protein	AB011166
high-mobility group (nonhistone chromosomal) protein 4	NM_005342
neuronal thread protein	NM_014486
GK001 protein	NM_020198
Heterogeneous nuclear ribonucleoprotein H1 (H)	AF753392
glutaminease	AF158555
Consensus Includes gb:AK028484.1 /DEF=Homo sapiens cDNA: FLJ22831 fls, clone KIAA1461, /FEA=mRNA /DB_XREF=gj:10439356 /UG=Hs.321666 Homo sapiens cDI	AK028484
pre-mRNA processing factor 31 homolog (yeast)	A1534332
mitochondrial ribosomal protein L19	NM_014763
macrophage stimulating, pseudogene 9	U28055
dopamine receptor D2	962137

double homeobox, 4	D38024	1.85
odd-skipped-related 2A protein	AI811238	1.85
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)	M58297	1.85
Interleukin 8 receptor, beta	NM_001557	1.85
G protein-coupled receptor 4	NM_005282	1.85
KIAA0007 protein	D26488	1.85
caspase 5, apoptosis-related cysteine protease	NM_004347	1.85
rab3 GTPase-activating protein, non-catalytic subunit (150kD)	AK021928	1.85
carboxypeptidase M	NM_001874	1.85
alkaline phosphatase, intestinal	NM_001631	1.84
myosin, heavy polypeptide 13, skeletal muscle	NM_003802	1.84
hypothetical protein PRO2533	NM_018629	1.84
decay accelerating factor for complement (CD55, Cromer blood group system)	BC001288	1.84
hypothetical protein PRO2800	NM_018635	1.84
glycophorin A (includes MN blood group)	U00170	1.84
Consensus includes gb:AK024136.1 /DEF=Homo sapiens cDNA FLJ14074 fs, clone HEMMB1001869, /FEA=mRNA /DB_XREF=gi:10436442 /UG=Hs.141208 Homo sapie	AK024136	1.84
hypothetical protein MGC3196	NM_024084	1.84
neuropeptide Y receptor Y6 (pseudogene)	U59431	1.84
Lsm1 protein	NM_014462	1.84
integrin, beta 5	AI335208	1.84
hypothetical protein FLJ23447	NM_024825	1.84
dipeptidase 1 (renal)	NM_004413	1.84
Consensus includes gb:AK025038.1 /DEF=Homo sapiens cDNA: FLJ21385 fs, clone COL03356, /FEA=mRNA /DB_XREF=gi:10437474 /UG=Hs.306750 Homo sapiens cC AK025038	AK025038	1.84
Consensus includes gb:AI570199 /FEA=EST /DB_XREF=gi:4533573 /DB_XREF=est:to76b07.x1 /CLONE=IMAGE:2184181 /UG=Hs.182183 Homo sapiens mRNA for cald, AI570199	AI570199	1.84
gb:BC000108.1 /DEF=Homo sapiens, Similar to Nedd-4-like ubiquitin-protein ligase, clone MGC:2079, mRNA, complete cds. /FEA=mRNA /PROD=Similar to Nedd-4-like u	BC000108	1.84
EphA4	BC004264	1.84
translation initiation factor IF2	BE138647	1.84
KIAA0138 gene product	D50928	1.83
immunoglobulin superfamily, member 4	AL519710	1.83
connective tissue growth factor	M92934	1.83
sex comb on midleg-like 2 (Drosophila)	NM_006089	1.83
Consensus includes gb:AK021571.1 /DEF=Homo sapiens cDNA FLJ11509 fs, clone HEMBA1002166, /FEA=mRNA /DB_XREF=gi:10432777 /UG=Hs.285093 Homo sapie	AK021571	1.83
IGF-II mRNA-binding protein 2	NM_006548	1.83
Consensus includes gb:AK027193.1 /DEF=Homo sapiens cDNA: FLJ23540 fs, clone LNG08239, /FEA=mRNA /DB_XREF=gi:10440282 /UG=Hs.293331 Homo sapiens cC AK027193	AK027193	1.83
eukaryotic translation initiation factor 4E	NM_001968	1.83
taste receptor, type 2, member 16	NM_016945	1.83
hypothetical protein FLJ10687	NM_018178	1.83
Consensus includes gb:AW301937 /FEA=EST /DB_XREF=gi:6711614 /DB_XREF=est:tr85b03.x1 /CLONE=IMAGE:2766893 /UG=Hs.138038 ESTs	AW301937	1.83
Consensus includes gb:BE786598 /FEA=EST /DB_XREF=gi:10207796 /DB_XREF=est:601478824F1 /CLONE=IMAGE:3881717 /UG=Hs.101474 KIAA0807 protein	BE786598	1.83
phospholipid scramblase 2	NM_020359	1.83
N-myc downstream-regulated gene 2	AI358939	1.83
SPARC-like 1 (masp8, hevjin)	NM_004684	1.83
RAB36, member RAS oncogene family	AF133588	1.83
Consensus includes gb:AK024103.1 /DEF=Homo sapiens cDNA FLJ14046 fs, clone HEMBA1006461, /FEA=mRNA /DB_XREF=gi:10436406 /UG=Hs.142677 Homo sapie	AK024108	1.83
TXK tyrosine kinase	AW276646	1.83
mitogen-activated protein kinase 8 interacting protein 2	NM_016431	1.83
RNA-binding protein gene with multiple splicing	DB4109	1.82
truncleotide repeat containing 15	AB014542	1.82
similar to rab11-binding protein	NM_019045	1.82
complement component 7	NM_000587	1.82
keratin 18	AL031903	1.82
hypothetical protein FLJ21625	NM_025039	1.82
pleckstrin homology-like domain, family A, member 1	A1759508	1.82
Consensus includes gb:AW973253 /FEA=EST /DB_XREF=gi:8163100 /DB_XREF=est:EST385351 /UG=Hs.292669 ESTs	AW973253	1.82
novel putative protein similar to YIL091C yeast hypothetical 84 kD protein from SGA1-KTR7	AI770084	1.82
hypothetical protein FLJ10486	BF591163	1.82
short stature homeobox 2	NM_006884	1.82
insulin	NM_000207	1.82
atractin	AL132773	1.81
retinoblastoma binding protein 6	NM_006910	1.81
Consensus includes gb:AL049260.1 /DEF=Homo sapiens mRNA: cDNA DXF2p564E233 (from clone DKF2p564E233), /FEA=mRNA /DB_XREF=gi:4500007 /UG=Hs.3020 /AL049260	AL049260	1.81
Motho	NM_004785	1.81
glycoprotein M6B	AI419030	1.81

Cdc42 guanine exchange factor (GEF) 9
 DKF ZP564J102 protein
 RAP78, member of RAS oncogene family
 CASP8 and FADD-like apoptosis regulator
 intermediate filament protein syncoilin
 hypothetical protein FLJ20048
 protein phosphatase 2 (formerly 2A), regulatory subunit B' (PR 72), alpha isoform and (PR 130), beta isoform
 MDS024 protein
 unc-51-like kinase 2 (C. elegans)
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)
 homeo box C8
 calcium/calmodulin-dependent serine protein kinase (MAGUK family)
 myosin, heavy polypeptide 1, skeletal muscle, adult
 HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)
 transducer of ERBB2, 2
 tyrosine aminotransferase
 sialophorin (gpL115, leukosialin, CD43)
 Consensus includes gb:AL050136.1 /DEF=Homo sapiens mRNA; cDNA DKF Zp586L141 (from clone DKF Zp586L141). /FEA=mRNA /DB_XREF=gi:4884346 /UG=Hs.14094
 sialic acid binding, Ig-like lectin 8
 tumor protein p53 (L-Fraumeni syndrome)
 oxidative stress induced like
 Consensus includes gb:BG280532 /FEA=EST /DB_XREF=gi:13047560 /DB_XREF=est:602388395F1 /CLONE=IMAGE:4517206 /UG=Hs.11210 ESTs, Moderately similar
 bromodomain, testis-specific
 prostatic binding protein
 alpha-2-glycoprotein 1, zinc
 erythrocyte membrane protein band 7.2 (stomatin)
 coagulation factor II (thrombin) receptor
 Consensus includes gb:AK026913.1 /DEF=Homo sapiens cDNA: FLJ23260 fls, clone COL05804, highly similar to HSU0911 Human clone 23652 mRNA sequence. /FEA=AK026913
 Consensus includes gb:AK026913.1 /DEF=Homo sapiens cDNA: FLJ23260 fls, clone COL05804, highly similar to HSU0911 Human clone 23652 mRNA sequence. /FEA=AK026913
 CLL7 protein
 neurotrophic tyrosine kinase, receptor, type 1
 serine protease inhibitor-like, with Kunitz and WAP domains 1 (epplin)
 L39424 B subtilis dapsB, kJf, kJf, kJf genes corresponding to nucleolides 1358-3197 of L39424 (-S, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
 zinc finger protein 161
 homolog of mouse BMP-2 inducible kinase
 COL142
 Consensus includes gb:A042030 /FEA=EST /DB_XREF=gi:3281224 /DB_XREF=est:ox60a09.x1 /CLONE=IMAGE:1660696 /UG=Hs.132153 ESTs
 U6 snRNA-associated Sm-like protein
 fatty acid desaturase 2
 actin related protein 2/3 complex, subunit 1B (41 kD)
 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C
 Consensus includes gb:BE326381 /FEA=EST /DB_XREF=gi:9200157 /DB_XREF=est:hw02a08.x1 /CLONE=IMAGE:3181766 /UG=Hs.18724 Homo sapiens mRNA; cDNA
 chromosome 21 open reading frame 82
 nudix (nucleoside diphosphate linked moiety X)-type motif 7
 Consensus includes gb:AL390026 /DEF=Human DNA sequence from clone RP3-336K20 on chromosome 6 Contains parts of 2 genes for novel proteins, ESTs, STSs and
 beta-amyloid binding protein precursor
 SMART/HDAC1 associated repressor protein
 transmembrane gamma-carboxyglutamic acid protein 4
 interleukin 4
 bromodomain adjacent to zinc finger domain, 1A
 hypothetical protein FLJ13769
 hypothetical protein FLJ10979
 Consensus includes gb:AK023017.1 /DEF=Homo sapiens cDNA FLJ12955 fls, clone NT2RP2005496, moderately similar to ZINC FINGER PROTEIN 135. /FEA=mRNA /DB_XREF=gi:9513716 /DB_XREF=est:hz76b04.x1 /CLONE=IMAGE:3213871 /UG=Hs.250175 homolog of yeast long chain
 v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)
 GTP binding protein 2
 mitogen-activated protein kinase kinase 5
 Consensus includes gb:BE467941 /FEA=EST /DB_XREF=gi:9513716 /DB_XREF=est:hz76b04.x1 /CLONE=IMAGE:3213871 /UG=Hs.250175 homolog of yeast long chain
 KIAA0186 gene product
 makorin, ring finger protein, 4
 testis specific ankyrin-like protein 1
 transcription elongation factor A (SII), 2

tyrosine kinase with immunoglobulin and epidermal growth factor homology domains
Williams-Beuren syndrome chromosome region 23
lactinin 1
SCAN domain-containing 2
v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog
T-box 2
N-myc downstream-regulated gene 2
p21 (CDKN1A)-activated kinase 2
HIC1-related gene on chromosome 22
hypothetical protein FLJ23516
ubiquitin-conjugating enzyme E2 variant 2
apolipoprotein F
Consensus includes gb:AW025578 /FEA=EST /DB_XREF=gi:5879109 /DB_XREF=est:uw75b04.x1 /CLONE=IMAGE:2525839 /UG=Hs.100261 Homo sapiens mRNA; cDNA, AW025578
azurocidin 1 (cationic antimicrobial protein 37)
adenomatous polyposis coli
Consensus includes gb:AA573805 /FEA=EST /DB_XREF=gi:2348320 /DB_XREF=est:nk07g11.s1 /CLONE=IMAGE:1012888 /UG=Hs.152939 Homo sapiens clone 24630 r AA573805
interleukin 18 (interferon-gamma-inducing factor)
hypothetical protein FLJ10701
hypothetical protein FLJ23392
CGI-148 protein
matrix metalloproteinase 10 (stromelysin 2)
DKFZP564A122 protein
CD47 antigen (Rb-related antigen, integrin-associated signal transducer)
Imprinted in Prader-Willi syndrome
Sm protein F
retinoblastoma-like 2 (p130)
signal sequence receptor, alpha (translocin-associated protein alpha)
5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)
myeloidlymphoid or mixed-lineage leukemia (t(11q24); translocated to, 10
hypothetical protein MGC2856
PTPRF interacting protein, binding protein 1 (liprin beta 1)
SmcX homolog, X chromosome (mouse)
interferon regulatory factor 2
KIAA0472 protein
phosphoenolpyruvate carboxykinase 1 (soluble)
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34
cyclic fibrosis transmembrane conductance regulator, ATP-binding cassette (sub-family C, member 7)
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily A, member 1
gb:NM_008273.2 /DEF=Homo sapiens small inducible cytokine A7 (monocyte chemoattractant protein 3) (SCYA7), mRNA, /FEA=mRNA /GEN=SCYA7 /PROD=monocyte chemokine mRNA; cDNA D4146852
hypothetical protein PRO2796
Consensus includes gb:N48852 /FEA=EST /DB_XREF=gi:1188018 /DB_XREF=est:y73d05.s1 /CLONE=IMAGE:279177 /UG=Hs.285625 Homo sapiens mRNA; cDNA D4146852
hypothetical protein FLJ14166
chymotrypsin C (caldecrin)
heat shock 70kD protein 8
receptor tyrosine kinase-like orphan receptor 1
acetylcholinesterase (YT blood group)
chromosome 6 open reading frame 32
mitogen-activated protein kinase kinase kinase 5
DKFZP566K023 protein
Consensus includes gb:AV753204 /FEA=EST /DB_XREF=gi:10911052 /DB_XREF=est:AV753204 /CLONE=NPD0B08 /UG=Hs.170287 ESTs
hypothetical protein FLJ2849 similar to signal peptidase SPC2223
KIAA1644 protein
T cell activation, increased late expression
KRA3 zinc finger protein KR18
gb:NM_024305.1 /DEF=Homo sapiens hypothetical protein MGC4278 (MGC4278), mRNA, /FEA=mRNA /GEN=MGC4278 /PROD=hypothetical protein MGC4278 /DB_XREF=NM_024305
neurofilament 3 (150kD medium)
T-box 5
AT2 receptor-interacting protein 1
Consensus includes gb:AK021967.1 /DEF=Homo sapiens cDNA FLJ11905 f1s, clone HEMBB1000050, /FEA=mRNA /DB_XREF=gi:10433276 /UG=Hs.273759 Homo sapiens cDNA AK021967
retinoid X receptor, alpha
cholesterol 25-hydroxylase
UDP-glucose dehydrogenase
KIAA0871 protein

144

gem (nuclear organelle) associated protein 4	NM_015487	1.73
melanoma antigen recognized by T cells 2	NM_018194	1.73
transcription factor (p38 interacting protein)	AW003334	1.73
CGI-58 protein	BC006637	1.73
gilia maturation factor, beta	NM_004124	1.73
serine/threonine kinase 6	NM_003158	1.73
JM27 protein	NM_007003	1.73
KIAA0874 protein	X80821	1.73
Consensus includes gb:AF339807.1 /DEF=Homo sapiens clone IMAGE:26186, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507345 /UG=Hs.137567 Homo sapiens c4 AF339807		1.73
D site of albumin promoter (albumin D-box) binding protein	D28468	1.73
lymphoid nuclear protein related to AF4	NM_002285	1.73
erythropoietin receptor	M34986	1.73
G protein-coupled receptor 44	NM_004778	1.73
tripartite motif-containing 37	AK022701	1.73
eukaryotic translation initiation factor 4 gamma, 1	BE966878	1.73
cyclin-dependent kinase 6	A1738463	1.73
Consensus includes gb:BG109855 /FEA=EST /DB_XREF=gi:12603361 /DB_XREF=est:50227952.1 /CLONE=IMAGE:4367306 /UG=Hs.49476 Homo sapiens done TUAJ BG109855		1.73
cadherin-like 22	AF035300	1.73
Consensus includes gb:AW173720 /FEA=EST /DB_XREF=gi:6439668 /DB_XREF=est:x1105.x1 /CLONE=IMAGE:2656929 /UG=Hs.271356 ESTs		1.73
Consensus includes gb:AL121900 /DEF=Human DNA sequence from clone RP11-379J5 on chromosome 20 Contains the last exon of the SEC23B gene for Sec23 (S. cert AL121900		1.73
spectrin, alpha, erythrocytic 1 (elliptocytosis 2)	W87601	1.72
small nuclear ribonucleoprotein polypeptide E	AF131850	1.72
etoposide-induced mRNA	AF131850	1.72
Consensus includes gb:BF693921 /FEA=EST /DB_XREF=gi:11979329 /DB_XREF=est:502082488.F1 /CLONE=IMAGE:4246917 /UG=Hs.180513 ATP-binding cassette, aut BF693921		1.72
Src-like adaptor	U44403	1.72
5' nucleotidase, mitochondrial	NM_020201	1.72
hypothetical protein FLJ10430	NM_018092	1.72
Consensus includes gb:AL137590.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K0810 (from clone DKFZp434K0610). /FEA=mRNA /DB_XREF=gi:5808330 /UG=Hs.30 AL137590		1.72
PERG11 family member in MHC class I region	NM_005844	1.72
GRO1 oncogene (melanoma growth stimulating activity, alpha)	NM_001511	1.72
Consensus includes gb:W86781 /FEA=EST /DB_XREF=gi:1400509 /DB_XREF=est:zh6a03.s1 /CLONE=IMAGE:416812 /UG=Hs.293736 ESTs		1.72
chorionic gonadotropin, beta polypeptide	W86781	1.72
CDC5 cell division cycle 5-like (S. pombe)	NM_000737	1.72
leucine-rich repeat-containing 5	AB007892	1.72
ubiquitin-like 4	NM_018103	1.72
myeloid lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4	BE543027	1.72
hairy homolog (Drosophila)	AL049698	1.72
zinc finger protein, subfamily 1A, 4 (Eos)	NM_005524	1.72
sodium bicarbonate transporter 4	NM_022465	1.72
choroideremia-like (Rab escort protein 2)	AF243499	1.72
Consensus includes gb:A419423 /FEA=EST /DB_XREF=gi:4265354 /DB_XREF=est:tt28d09.x1 /CLONE=IMAGE:2097521 /UG=Hs.172780 ESTs		1.72
lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)	NM_001821	1.72
Consensus includes gb:A1865582 /FEA=EST /DB_XREF=gi:4897876 /DB_XREF=est:td8e05.x1 /CLONE=IMAGE:2268608 /UG=Hs.283824 Homo sapiens Skn-1a1-like mF A1865582		1.72
Consensus includes gb:AW268081 /FEA=EST /DB_XREF=gi:565511 /DB_XREF=est:rs2e03.x1 /CLONE=IMAGE:2767612 /UG=Hs.193408 ESTs, Weakly similar to ALU AW268081		1.72
one out domain, family member 1	U96173	1.72
hypothetical protein FLJ13885	NM_025016	1.72
gb:NM_018541.1 /DEF=Homo sapiens hypothetical protein PRO2832 (PRO2832), mRNA. /FEA=mRNA /GEN=PRO2832 /PROD=hypothetical protein PRO2832 /DB_XREF=gi:1018541		1.72
KIAA0328 protein	AB002328	1.72
caveolin 2	AA150110	1.72
phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma)	NM_005084	1.72
apolipoprotein C1	W76394	1.72
pleckstrin 2 (mouse) homolog	NM_016445	1.72
hypothetical protein similar to rat vacuolar sorting protein 33a	NM_022916	1.71
cytochrome P450, subfamily X1B (steroid 11-beta-hydroxylase), polypeptide 2	X54741	1.71
complement factor H-related 4	NM_006684	1.71
Consensus includes gb:AF023203.1 /DEF=Homo sapiens homeobox protein Ogl12 (OGL12) mRNA, partial cds. /FEA=mRNA /GEN=OGL12 /PROD=homeobox protein Ogl12 /AF023203		1.71
Max-interacting protein	BE502432	1.71
chromosome 5 open reading frame 7	NM_016604	1.71
rat guanine nucleotide dissociation stimulator	A421559	1.71
melanoma antigen, family D, 2	A1924630	1.71
variable charge, Y chromosome	NM_004879	1.71
H2A histone family, member O	BC001629	1.71
forkhead box G1A	NM_004471	1.71

KIAA1036 protein	NM_014909	1.71
homolog of mouse BMP-2 inducible kinase	AB015331	1.71
CD34 antigen	M61104	1.71
deiodinase, iodothyronine, type I	NM_000792	1.71
ribosome binding protein 1 homolog 180kD (dog)	AA706065	1.71
hypothetical protein FLJ10201	NM_018023	1.71
bridging integrator 2	NM_016293	1.71
mitogen-activated protein kinase kinase 11	NM_002419	1.71
hypothetical protein PRO1496	NM_018603	1.71
integrin, alpha L (anigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide)	AC002310	1.71
hypothetical protein FLJ10871	NM_018250	1.71
nuclear receptor binding factor-2	AA683074	1.71
retinoic acid repressible protein	AF172066	1.71
decorin	AF138303	1.71
RAD1 homolog (S. pombe)	NM_002853	1.71
glutamate receptor, metabotropic 7	NM_000844	1.71
Consensus includes gb:AF070547.1 /DEF=Homo sapiens clone 24820 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387909 /UG=Hs.146312 Homo sapiens clone 24820 AF070547	AF081459	1.70
origin recognition complex, subunit 5-like (yeast)	AF035594	1.70
Consensus includes gb:AF035594.1 /DEF=Homo sapiens protein kinase C-alpha mRNA, partial 3 UTR. /FEA=mRNA /DB_XREF=gi:3168857 /UG=Hs.279856 Homo sapiens clone 24820 AF070547	NM_021183	1.70
hypothetical protein similar to small G proteins, especially RAP-2A	NM_016242	1.70
endomucin-2	NM_001397	1.70
endothelin converting enzyme 1	AF260113	1.70
cytochrome P450 polypeptide 43	NM_014936	1.70
ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function)	AB001328	1.70
soluble carrier family 15 (oligopeptide transporter), member 1	NM_008865	1.70
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 3	AA634446	1.70
Consensus includes gb:AA634446 /FEA=EST /DB_XREF=gi:2557680 /DB_XREF=est:79e01.s1 /CLONE=IMAGE:744216 /UG=Hs.116840 ESTs	NM_014498	1.70
golgi phosphoprotein 4	NM_020157	1.70
coregulin	AF141392	1.70
RAN binding protein 7	AW238632	1.70
KIAA1098 protein	AK026548	1.70
KIAA0088 protein	AK025552	1.70
Consensus includes gb:AK025552.1 /DEF=Homo sapiens cDNA: FLJ21895 fis, clone HEP03467. /FEA=mRNA /DB_XREF=gi:10438101 /UG=Hs.306815 Homo sapiens clone 24820 AF070547	AK026120	1.70
KIAA0460 protein	AC005255	1.70
olfactory receptor, family 7, subfamily A, member 17	Y15014	1.70
UDP-Gal:beta-GlcNAc beta 1,3-galactosyltransferase, polypeptide 2	BF438282	1.70
POZ domain containing guanine nucleotide exchange factor(GEF)1	BF058643	1.70
EGF-like repeats and discoidin-like domains 3	NM_000726	1.70
calcium channel, voltage-dependent, beta 4 subunit	M80927	1.70
chitinase 3-like 1 (cartilage glycoprotein-39)	M24900	1.70
nuclear receptor subfamily 1, group D, member 1	BC003541	1.70
hypothetical protein FLJ10488	AL049215	1.70
Dystonia musculorum of mouse, human homolog of	L25275	1.70
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3	NM_017488	1.70
adducin 2 (beta)	AB028816	1.70
phosphodiesterase 10A	R85779	1.69
spectrin, beta, non-erythrocytic 1	AL049394	1.69
hypothetical protein FLJ10648	NM_004108	1.69
ficollin (collagen/vibroninogen domain-containing lectin) 2 (hu collagen)	AW301305	1.69
KIAA1096 protein	AB08958	1.69
KIAA0870 protein	NM_031275	1.69
testis expressed sequence 12	AL022185	1.69
carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7	AL137592	1.69
dual oxidase 1	NM_025057	1.69
hypothetical protein FLJ23189	NM_023613	1.69
KIAA1009 protein	AW974823	1.69
Consensus includes gb:AW974823 /FEA=EST /DB_XREF=gi:8166026 /DB_XREF=est:EST386928 /UG=Hs.105411 ESTs	NM_003555	1.69
olfactory receptor, family 1, subfamily G, member 1	NM_004407	1.69
dentin matrix acidic phosphoprotein	BE563442	1.69
interleukin 1 receptor antagonist	NM_014289	1.69
calpain 6	AL134489	1.69
ribonuclease P (38kD)	X56687	1.69
upstream binding transcription factor, RNA polymerase I		

147

slatytransferase 8 (alpha-2, 8-sialyltransferase) B	NM_006011	1.66
melanoma cell adhesion molecule	M29277	1.66
fibroblast growth factor 17	NM_003887	1.66
solute carrier family 1 (glial high affinity glutamate transporter), member 2	NM_004171	1.66
KIAA0683 gene product	AL137394	1.67
HSKM-B protein	AB09870	1.67
M11187 Human 28S rRNA sequence, length 5025 bases, 3 prime target bases 3331-4985	M11187	1.67
hydridiform mole associated and imprinted	AF241534	1.67
zinc finger protein 193	NM_006299	1.67
Consensus includes gb:AK022045.1 /DEF=Homo sapiens cDNA FLJ11983 fs, clone HEMBB1001337 /FEA=mRNA /DB_XREF=gi:10433364 /UG=Hs.193162 Homo sapiens mRNA; cDNA AK022045	NM_006299	1.67
Consensus includes gb:AW173157 /FEA=EST /DB_XREF=gi:6439105 /DB_XREF=est:84812.x1 /CLONE=IMAGE:2663502 /UG=Hs.26860 Homo sapiens mRNA; cDNA AW173157	AW173157	1.67
glycerol-3-phosphate dehydrogenase 1 (soluble)	AI368018	1.67
hemoglobin, gamma G	NM_000184	1.67
villigo-associated protein VIT-1	NM_018693	1.67
protein kinase, interferon-inducible double stranded RNA dependent	NM_002759	1.67
zinc finger protein 223	NM_013381	1.67
hypothetical protein PRO2714	NM_018534	1.67
aldehyde oxidase 1	AB046692	1.67
hepatocyte nuclear factor 4, alpha	Z49825	1.67
MAD, mothers against decapentaplegic homolog 5 (Drosophila)	AF010601	1.67
hypothetical protein MGC2508	AA639585	1.67
dyslin, cytoplasmic, intermediate polypeptide 2	AF250307	1.67
insulin-like growth factor binding protein 2 (38kD)	NM_000597	1.67
hypocretin (orexin) receptor 2	NM_001526	1.67
hypothetical protein FLJ22782	NM_024821	1.67
disca, large (Drosophila) homolog 1	BC251175	1.67
beta-site APP-cleaving enzyme	NM_012104	1.67
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD	NM_034777	1.67
v-akt murine thymoma viral oncogene homolog 1	NM_005163	1.67
transcription factor (p38 interacting protein)	NM_000861	1.67
transcription factor (p38 interacting protein)	NM_017569	1.67
topoisomerase (DNA) I	AW025108	1.67
hypothetical protein FLJ13340	NM_025085	1.67
RNA binding motif protein 12	AL514547	1.67
placental growth factor, vascular endothelial growth factor-related protein	BC001422	1.67
EST-YD1 protein	NM_021208	1.67
putative X-linked retinopathy protein	SS8722	1.67
lethal (3) malignant brain tumor (3)mbt protein (Drosophila) homolog	AW445040	1.67
aquaporin 1 (channel-forming integral protein, 28kD)	AL518391	1.67
KIAA0841 protein	AA845355	1.67
hypothetical gene CG018	AL049786	1.67
lysosomal-associated membrane protein 3	NM_014398	1.67
solute carrier family 1 (glutamate transporter), member 7	NM_006671	1.67
KIAA0908 protein	AI867102	1.66
aldehyde dehydrogenase 3 family, member B2	NM_000685	1.66
protease, serine, 12 (neutrypsin, molopsin)	AI810787	1.66
KIAA0528 gene product	AB011100	1.66
KIAA0675 gene product	NM_014648	1.66
manic fringe homolog (Drosophila)	AI780053	1.66
retinoic acid receptor, beta	X04014	1.66
basement membrane-induced gene	AB035482	1.66
von Hippel-Lindau syndrome	NM_000551	1.66
general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD)	NM_001521	1.66
insulin-like growth factor binding protein 5	M65062	1.66
interleukin 1, zeta	NM_014439	1.66
cytochrome P450, subfamily IIA (riphedipine oxidase), polypeptide 4	NM_017460	1.66
Consensus includes gb:AB007970.1 /DEF=Homo sapiens mRNA, chromosome 1 specific transcript KIAA0501, /FEA=mRNA /DB_XREF=gi:3413945 /UG=Hs.223020 Hom	AB007970	1.66
chymotrypsinogen B1	NM_001906	1.66
Sp3 transcription factor	AI145005	1.66
gamma-glutamyl carboxylase	BE326952	1.66
Consensus includes gb:AK024553.1 /DEF=Homo sapiens cDNA: FLJ20900 fs, clone ADKA03593, /FEA=mRNA /DB_XREF=gi:10436861 /UG=Hs.306887 Homo sapiens c	AK024553	1.66
Consensus includes gb:AK025072.1 /DEF=Homo sapiens cDNA: FLJ21419 fs, clone COL04084, /FEA=mRNA /DB_XREF=gi:10437512 /UG=Hs.306759 Homo sapiens c	AK025072	1.66
zinc finger protein 238	AK000847	1.66

hypothetical protein FLJ13505 NM_024577 1.66
 Consensus includes gb:AU145680 /FEA=EST /DB_XREF=gi:1007201 /DB_XREF=est:AU145680 /CLONE=HEMBA1005500 /UG=Hs.306613 Homo sapiens cDNA FLJ11: AU145680 1.66
 hypothetical protein MGC5528 L11372 1.66
 hypothetical protein KIAA1102 protein AK028815 1.66
 kynureninase (L-kynurenine hydrolase) NM_003937 1.66
 muscle-specific beta 1 integrin binding protein NM_014446 1.66
 chromosome 21 open reading frame 2 NM_004928 1.66
 hypothetical protein PRO1584 NM_018586 1.66
 neuromedin B receptor NM_002511 1.66
 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e NM_007100 1.66
 actin-like complex-like 1 (Drosophila) BC001638 1.66
 protein phosphatase 2, regulatory subunit B (B56), alpha isoform AB03525 1.66
 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47) M31523 1.66
 KIAA0680 gene product AW295193 1.66
 KIAA0414 protein T90308 1.66
 Consensus includes gb:AW299245 /FEA=EST /DB_XREF=gi:6708922 /DB_XREF=est:xs48d07.x1 /CLONE=IMAGE:2772877 /UG=Hs.62601 Homo sapiens mRNA: cDNA 1.66
 melastase chromosome protein 1 NM_017513 1.66
 protocadherin alpha 9 NM_014005 1.66
 hypothetical protein FLJ10044 NM_017980 1.66
 potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 NM_002252 1.66
 transcription factor 7 (T-cell specific, HMG-box) NM_003202 1.65
 mitogen-activated protein kinase 8 interacting protein 2 NM_012324 1.65
 chemokine (C-C motif) receptor 2 NM_000648 1.65
 Consensus includes gb:BF434424 /FEA=EST /DB_XREF=gi:11446712 /DB_XREF=est:768608.x1 /CLONE=IMAGE:3844393 /UG=Hs.107184 spectrin, beta, non-erythrocyte BF434424 1.65
 Consensus includes gb:AU147942 /FEA=EST /DB_XREF=gi:1009463 /DB_XREF=est:AU147942 /CLONE=MAMMA1002188 /UG=Hs.287517 Homo sapiens cDNA FLJ12 AU147942 1.65
 olfactory receptor, family 10, subfamily H, member 1 NM_013940 1.65
 Consensus includes gb:BC004145.1 /DEF=Homo sapiens, clone IMAGE:3509783, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3509783) /DB_XI BC004145 1.65
 serine/threonine kinase 11 (Peutz-Jeghers syndrome) NM_000455 1.65
 RAB31, member RAS oncogene family BE789881 1.65
 chromosome 2 open reading frame 3 BC000853 1.65
 thyrotropin-releasing hormone degrading ectoenzyme NM_013381 1.65
 pavin, alpha NM_018222 1.65
 tumor endothelial marker 7 precursor AF070526 1.65
 flavin containing monooxygenase 2 BC005894 1.65
 gastrulation brain homeo box 2 AF118452 1.65
 thrombospondin 1 NM_003246 1.65
 Ccort10 protein AF214738 1.65
 hypothetical protein FLJ12121 NM_024978 1.65
 tumor necrosis factor (ligand) superfamily, member 13 AF114012 1.65
 KIAA0339 gene product N30342 1.65
 Cluster incl. AL050186 Homo sapiens mRNA: cDNA DKFZp586D1122 (from clone DKFZp586D1122) /cds=UNKNOWN /gb=AL050166 /ig=4884381 /ug=Hs.26295 /len=26: 1.65
 MAX binding protein NM_020310 1.65
 integrin, alpha 2b (platelet glycoprotein IIb of fibrin complex, antigen CD41B) J02863 1.65
 SREBP CLEAVAGE-ACTIVATING PROTEIN D83782 1.65
 inhibitor, beta B (ectonin AB beta polypeptide) NM_002193 1.65
 O-substrate NM_006658 1.65
 LIM domain protein AF153882 1.65
 enhancer of zeste homolog 1 (Drosophila) AB004818 1.65
 glutamate receptor, metabotropic 1 NM_000838 1.65
 AND-1 protein AK001538 1.65
 microtubule-associated protein 1A AA633827 1.65
 arachidonate 12-lipoxygenase NM_000697 1.65
 purine-rich element binding protein G NM_013357 1.65
 lamin A/C AA063189 1.65
 thyroid hormone receptor interactor 6 AF000874 1.65
 KIAA0870 protein AB020877 1.64
 myelin-associated oligodendrocyte basic protein D28114 1.64
 Consensus includes gb:AL117643.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434M245 (from clone DKFZp434M245) /FEA=mRNA /DB_XREF=gi:5912233 /UG=Hs.5286 AL117643 1.64
 BAI1-associated protein 3 NM_004827 1.64
 thyroid transcription factor 1 BC006221 1.64
 hypothetical protein PRO2007 NM_018613 1.64
 immunoglobulin heavy constant gamma 3 (G3m marker) AJ275469 1.64

transferrin
Consensus includes gb:AU147962 /FEA=EST /DB_XREF=gi:11009483 /DB_XREF=est:AU147962 /CLONE=MAMMA1002250 /UG=Hs.301509 Homo sapiens cDNA FLJ12
NM_006663 1.64
RelA-associated inhibitor
AL078459 1.64
dimethylarginine dimethylaminohydrolase 1
AW182367 1.64
metal-regulatory transcription factor 1
L32610 1.64
heterogeneous nuclear ribonucleoprotein H3 (2H9)
NM_015848 1.64
cyclotriptin 2
NM_015848 1.64
solute carrier family 28 (sulfate transporter), member 2
AI025519 1.64
adaptor-related protein complex 1, sigma 1 subunit
BC003561 1.64
phospholipase A2 receptor 1, 180kD
U17033 1.64
ACO for serine protease homologue
NM_017509 1.64
hypothetical protein FLJ12688
NM_024997 1.64
Consensus includes gb:W28849 /FEA=EST /DB_XREF=gi:1308815 /DB_XREF=est:52g6 /UG=Hs.33862 ESTs
W28849 1.64
PRO0097 protein
NM_014114 1.64
HCNP protein; XPA-binding protein 2
NM_020196 1.64
neurophilin 3
AI837333 1.64
solute carrier family 6 (neurotransmitter transporter, laurine), member 6
BC006252 1.64
BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like
AF288153 1.64
Consensus includes gb:AL121903 /DEF=Human DNA sequence from clone RP1-155G6 on chromosome 20 Contains part of the gene for brefeldin A-inhibited guanine nud
AL121903 1.64
BH-protocadherin (brain-heart)
AB006757 1.64
HYA22 protein
BF002474 1.64
zinc finger protein 175
NM_007147 1.64
F-box only protein 9
AL137520 1.64
CAMP response element-binding protein CRE-BPa
NM_004904 1.64
NBR2
NM_005821 1.64
ATP-binding cassette, sub-family A (ABC1), member 1
NM_005502 1.64
SON DNA binding protein
AI836458 1.64
Phosphatidylylcytophosphate Synthase
NM_024419 1.64
CGI-43 protein
NM_015622 1.64
Escherichia coli /REF=J04423 /DEF=E coli bioD gene dehiobiotin synthetase corresponding to nucleotides 5024-5244 of J04423 /LEN=876 (-5 and -3 represent transcript r
J04423 1.64
tachykinin receptor 1
M81797 1.64
activating transcription factor 1
AA34345 1.64
Consensus includes gb:AL078638.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIIMAGE 117929. /FEA=mRNA /DB_XREF=gi:5051866 /UG=Hs.227171 AL078638
AL078638 1.63
S100 calcium binding protein, beta (neural)
BC001766 1.63
pancreatic polypeptide 2
NM_021092 1.63
Consensus includes gb:AU147200 /FEA=EST /DB_XREF=gi:11008721 /DB_XREF=est:AU147200 /CLONE=HEMBB1002699 /UG=Hs.296698 Homo sapiens cDNA FLJ12
AU147200 1.63
homonally upregulated Neu-associated kinase
NM_014586 1.63
guanine nucleotide binding protein (G protein), beta 5
NM_016194 1.63
F-box only protein 22
NM_012170 1.63
Consensus includes gb:AF070602.1 /DEF=Homo sapiens cDNA 24504 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387982 /UG=Hs.51649 Homo sapiens cDNA 24504 /AF070602
AF070602 1.63
Consensus includes gb:AK022888.1 /DEF=Homo sapiens cDNA FLJ2624 fls. clone NT2RM4001754. /FEA=mRNA /DB_XREF=gi:10434225 /UG=Hs.306639 Homo sapien
AK022888 1.63
HIV-1 inducer of short transcripts binding protein; lymphoma related factor
AW027070 1.63
mitochondrial solute carrier
NM_018579 1.63
collagen, type VIII, alpha 1
BE877796 1.63
likely homolog of rat and mouse retinoid-inducible serine carboxypeptidase
NM_021626 1.63
nuclear receptor coactivator 1
AI872428 1.63
WW domain binding protein 4 (formin binding protein 21)
NM_007187 1.63
gb:NM_018834.1 /DEF=Homo sapiens hypothetical protein PRO2893 (PRO2893), mRNA. /FEA=mRNA /GEN=PRO2893 /PROD=hypothetical protein PRO2893 /DB_XREF
NM_018834 1.63
ribosomal protein S8 kinase, 70kD, polypeptide 2
NM_003592 1.63
muscle, skeletal, receptor tyrosine kinase
BE222739 1.63
purine-rich element binding protein A
NM_004235 1.63
Kruppel-like factor 4 (gut)
NM_000377 1.63
Wiskott-Aldrich syndrome (eczema-thrombocytopenia)
NM_003888 1.63
aldehyde dehydrogenase 1 family, member A2
Consensus includes gb:AL108696.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIIMAGE 21920. /FEA=mRNA /DB_XREF=gi:5689771 /UG=Hs.185701 /AL108696
AL108696 1.63
Consensus includes gb:AW451230 /FEA=EST /DB_XREF=gi:8992006 /DB_XREF=est:UH-H-B13-ah-a-01-Q-U1.s1 /CLONE=IMAGE:2736649 /UG=Hs.257332 ESTs. Highly: AW451230
AW451230 1.63
fibronectin leucine rich transmembrane protein 3
NM_021281 1.63
phosphotyrosine phosphohistidine inorganic pyrophosphate phosphatase
NM_022126 1.63
chondroin sulfate proteoglycan 4 (melanoma-associated)
NM_001897 1.63
arrestin A2 pseudogene 1
M82896 1.63
Consensus includes gb:AL121881 /DEF=Human DNA sequence from clone RPS-1061C18 on chromosome 1p36.22-36.33. Contains ESTs, STSs, GSSs and a CoQ Island. AL121881
AL121881 1.63
sterile-alpha motif and leucine zipper containing kinase AZK
NM_016653 1.63
DKF ZP586J0119 protein
AF112207 1.63

KIAA0138 gene product 1.61
aldehyde dehydrogenase 1 family, member A2 NM_014649
heat shock 70kD protein 2 1.61
hypothetical gene FLJ00060 AG013226
7-dehydrocholesterol reductase 1.61
leucine zipper, down-regulated in cancer 1 AK024467
karyopherin beta 2b, transportin NM_001360
WW domain-containing oxidoreductase 1.61
cell division cycle 25B AK022217
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson relaxin 2 (H2) 1.61
synaptotagmin XIII AL109804
hypothetical protein My014 NM_022969
like mouse brain protein E46 NM_005059
putative nuclear protein AW099736
mitogen-activated protein kinase kinase 3 NM_030918
ubiquitin-conjugating enzyme E2B (RAD6 homolog) Z95331
Consensus includes gb:AF052160.1 /DEF=Homo sapiens clone 24629 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360471 /UG=Hs.142570 Homo sapiens clone 24629 AF052160
angiotensin-like 2 NM_012098
SEC22 vesicle trafficking protein-like 1 (S. cerevisiae) 1.61
tumor necrosis factor (ligand) superfamily, member 13 1.61
J04423 E coli bioD gene deshydrobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively) 1.61
ATPass, Ca++ transporting, plasma membrane 2 AF114013
hypothetical protein MGC2742 J04423
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A) R52647
fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor) NM_023938
Consensus includes gb:AF319523.1 /DEF=Homo sapiens RT-LI mRNA, complete sequence. /FEA=mRNA /PROD=RT-LI /DB_XREF=gi:13310144 /UG=Hs.326775 Homo sapiens AF319523
UDP-Gal-4-epiChNAc beta 1,3-galactosyltransferase, polypeptide 5 NM_008057
Consensus includes gb:X61070.1 /DEF=Human mRNA for T cell receptor, clone IGRA15. /FEA=mRNA /PROD=T cell receptor /DB_XREF=gi:333508 /UG=Hs.121492 Homo sapiens X61070
Interleukin 1 receptor accessory protein AF167343
Consensus includes gb:AW293343 /FEA=EST /DB_XREF=gi:6699905 /DB_XREF=est:U1-H-B12-ah1-d-05-Q-U1.s1 /CLONE=IMAGE:2726841 /UG=Hs.283978 Homo sapiens AW293343
LIM domain kinase 1 NM_002314
ATP-binding cassette, sub-family B (MOTAP), member 11 NM_003742
dipeptidylpeptidase IV (CD26, adenosine deaminase complexing protein 2) M80536
musculin (activated B-cell factor-1) 1.61
hypothetical protein MGC5489 AF060154
KIAA0042 gene product NM_024055
KIAA1334 protein NM_014875
CGI-148 protein A4460694
ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome) NM_016078
Consensus includes gb:AL096727.1 /DEF=Homo sapiens mRNA: cDNA DKFZP434B104 (from clone DKFZP434B104). /FEA=mRNA /DB_XREF=gi:5419860 /UG=Hs.1682 AL096727
aldehyde dehydrogenase 7 family, member A1 BC002515
APMCF1 protein NM_021203
purinergic receptor P2X, ligand-gated ion channel, 4 NM_002560
Consensus includes gb:AK023174.1 /DEF=Homo sapiens cDNA: FLJ11521 fs, clone COL05880. /FEA=mRNA /DB_XREF=gi:10437836 /UG=Hs.306777 Homo sapiens cDNA: AK023174
zinc finger protein 284 AL163248
hyaluronoglucosaminidase 4 NM_012269
carbonic anhydrase III, muscle specific NM_005181
hypothetical protein FLJ20783 W87468
carbonyl reductase 1 BC002511
gamma-glutamyltransferase 2 A344075
ankyrin 2, neuronal NM_001148
hypothetical protein MGC3038 NM_023942
potassium inwardly-rectifying channel, subfamily J, member 15 U73191
cardiotrophin 1 NM_001330
calcitonin/calcitonin-related polypeptide, alpha M28095
Consensus includes gb:BF197222 /FEA=EST /DB_XREF=gi:11085906 /DB_XREF=est:7m8807.x1 /CLONE=IMAGE:3561949 /UG=Hs.12342 Homo sapiens clone 24538 BF197222
CD5 antigen-like (scavenger receptor cysteine rich family) 1.60
CGI-109 protein NM_005894
cyclochrome P450, subfamily VIIIB (cytochrome P450, subfamily VIIIB (alpha-hydroxylase), polypeptide 1 AF151867
Consensus includes gb:AK001448.1 /DEF=Homo sapiens cDNA FLJ10588 fs, clone NT2RP2003986. /FEA=mRNA /DB_XREF=gi:7022713 /UG=Hs.301255 Homo sapiens AK001448
glutamate receptor, metabotropic 5 NM_000842
preproelastin-releasing peptide NM_015893

ectopic viral integration site 5
 N-acylsphingosine amidohydrolase (acid ceramidase)
 Consensus includes gb:BG230612 /FEA=EST /DB_XREF=gi:12725654 /DB_XREF=estn41b09.x1 /CLONE=IMAGE:4143329 /UG=Hs.13998 Homo sapiens cDNA: FLJ22: BG230612
 RBP1-like protein
 signal transducing adaptor molecule (SH3 domain and ITAM motif) 2
 hypothetical protein FLJ20856
 CED-6 protein
 zinc finger protein 76 (expressed in testis)
 cellular retinoic acid binding protein 2
 Consensus includes gb:U78287.1 /DEF=Human clone 23589 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710280 /UG=Hs.11506 Human clone 23589 mRNA sequence U79297
 Consensus includes gb:AA700015 /FEA=EST /DB_XREF=gi:2702978 /DB_XREF=estz69c11.s1 /CLONE=IMAGE:436052 /UG=Hs.287995 Homo sapiens cDNA: FLJ2318 AA700015
 GTP binding protein overexpressed in skeletal muscle
 KIAA0416 protein
 thrombospondin 2
 growth arrest-specific 2
 Consensus includes gb:AA653456 /FEA=EST /DB_XREF=gi:2589627 /DB_XREF=estag66c08.s1 /CLONE=IMAGE:1127918 /UG=Hs.270495 ESTs, Weakly similar to ubiq
 solute carrier family 2 (facilitated glucose transporter), member 3
 Cyclin E1
 KIAA1451 protein
 Microfibril-associated glycoprotein-2
 sodium channel, voltage-gated, type IV, alpha polypeptide
 Consensus includes gb:R78289 /FEA=EST /DB_XREF=gi:853409 /DB_XREF=estiy81n12.s1 /CLONE=IMAGE:145703 /UG=Hs.287419 Homo sapiens cDNA: FLJ11505 fts R78289
 hypothetical protein FLJ12681
 apolipoprotein A-II
 perlecan
 KIAA0987 protein
 hypothetical protein PRO2435
 Consensus includes gb:AA491625 /FEA=EST /DB_XREF=gi:2221187 /DB_XREF=estne80d01.s1 /CLONE=IMAGE:910561 /UG=Hs.283940 ESTs
 S100 calcium binding protein A12 (calgranulin C)
 keratin 17
 hemochromatosis
 annexin A5
 synaptobrevin 1
 chromosome 8 open reading frame 10
 basic leucine zipper nuclear factor 1 (JEM-1)
 activating transcription factor 5
 gb:BC006361.1 /DEF=Homo sapiens, clone MGC:13137, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13137) /DB_XREF=gi:13823508 /FL=gb:BC006361
 gb:U35622.2 /DEF=Homo sapiens EWS proteinE1A enhancer binding protein chimera mRNA, complete cds. /FEA=mRNA /PROD=EWS proteinE1A enhancer binding prot U35622
 retinoblastoma binding protein 1
 solute carrier family 4, sodium bicarbonate cotransporter, member 4
 estrogen receptor 1
 hypothetical protein FLJ10461
 neighbor of A-kinase anchoring protein 95
 FYN binding protein (FYB-120/130)
 hypothetical protein MGC5306
 skin-specific protein
 gb:AF202635.1 /DEF=Homo sapiens PP1200 mRNA, complete cds. /FEA=mRNA /PROD=PP1200 /DB_XREF=gi:10732645 /UG=Hs.302135 Homo sapiens PP1200 mRNA AF202635
 hypothetical protein PRO1438
 G protein-coupled receptor
 A kinase (PRKA) anchor protein 6
 GDNF family receptor alpha 1
 amyloid beta (A4) precursor-like protein 2
 fibroblast growth factor 12
 hypothetical protein DKFZp761G1913
 KIAA0346 protein
 vav 2 oncogene
 hypothetical protein FLJ20004
 2.19 gene
 Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2
 aggrecan 1 (chondroitin sulfate proteoglycan, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122)
 formyl peptide receptor 1
 KIAA1025 protein
 KIAA0669 gene product

1.60 NM_005665
 1.60 U4764
 1.60 BG230612
 1.60 AL133600
 1.60 NM_025143
 1.60 AK023668
 1.60 NM_003427
 1.60 NM_001878
 1.60 U79297
 1.60 AA700015
 1.60 NM_005261
 1.60 NM_015564
 1.60 NM_003247
 1.60 NM_005256
 1.60 AA653456
 1.60 NM_006931
 1.60 AI671049
 1.60 BF970829
 1.60 AW665892
 1.60 NM_000334
 1.60 R78289
 1.60 NM_022773
 1.60 NM_001643
 1.60 AB046840
 1.60 NM_014950
 1.60 NM_018527
 1.60 AA491625
 1.60 NM_005621
 1.60 Z19574
 1.60 AF150864
 1.60 NM_001154
 1.60 NM_003895
 1.60 NM_008781
 1.60 NM_003666
 1.60 AF101388
 1.60 BC006361
 1.60 U35622
 1.60 NM_002892
 1.60 AF011390
 1.59 AI073549
 1.59 NM_018098
 1.59 AB016332
 1.59 AF198052
 1.59 NM_024116
 1.59 NM_014357
 1.59 AF202635
 1.59 NM_018600
 1.59 U73531
 1.59 NM_004274
 1.59 NM_005264
 1.59 BC000373
 1.59 AL119322
 1.59 AL136557
 1.59 AA521267
 1.59 NM_003371
 1.59 NM_017616
 1.59 BC002934
 1.59 AF109161
 1.59 NM_013227
 1.59 NM_002029
 1.59 BG426889
 1.59 AF201291

hypothetical protein FLJ10142	1.59	NM_018008
putative acid-sensing ion channel	1.59	NM_018674
Consensus includes gb:AL050198.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586D0923 (from clone DKFZp586D0923). /FEA=mRNA /DB_XREF=gi:4894439 /UG=Hs.62	1.59	AL050198
Consensus includes gb:AK022387.1 /DEF=Homo sapiens cDNA FLJ12335 fls, clone MAMMA1002219, highly similar to Rattus norvegicus rexo70 mRNA. /FEA=mRNA /DB_XREF=gi:4894439	1.59	AK022387
type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	1.59	AA195244
Consensus includes gb:AK025388.1 /DEF=Homo sapiens cDNA: FLJ21735 fls, clone COLF3350. /FEA=mRNA /DB_XREF=gi:10437892 /UG=Hs.306808	1.59	AK025388
hypothetical protein FLJ22215	1.59	NM_022834
homeodomain interacting protein kinase 2	1.59	NM_022740
endosulfine alpha	1.59	AF067170
PRKC, apoptosis, WT1, regulator	1.59	BF732462
DEAD/HD (Asp-Glu-Ala-Asp/His) box polypeptide 3	1.59	NM_001356
GRO2 oncogene	1.59	M57731
protein kinase, cAMP-dependent, catalytic, gamma	1.59	NM_002732
SEC14-like 2 (S. cerevisiae)	1.59	NM_012428
RAR-related orphan receptor A	1.59	U04897
prothymosin a14	1.59	NM_016171
empty spiracles homolog 1 (Drosophila)	1.59	AW196403
SRF (sex determining region Y)-box 20	1.59	AB023535
Consensus includes gb:AA555096 /FEA=EST /DB_XREF=gi:2325635 /DB_XREF=est:NM040e01.x1 /CLONE=IMAGE:594968 /UG=Hs.26295	1.59	AA555096
customer protein complex, subunit beta	1.59	NM_016451
carboxylesterase 2 (intestine, liver)	1.59	BF033242
sucrase-isomaltase	1.59	NM_001041
SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	1.59	BC001364
Consensus includes gb:AK025380.1 /DEF=Homo sapiens cDNA: FLJ21707 fls, clone COL09953. /FEA=mRNA /DB_XREF=gi:10437861 /UG=Hs.306808	1.59	AK025380
surfactant, pulmonary-associated protein D	1.59	NM_003019
POM (POM121 rat homolog) and ZP3 fusion	1.59	AB344473
deca, large (Drosophila) homolog 5	1.59	NM_012230
phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	1.59	AB011155
DEAD/HD (Asp-Glu-Ala-Asp/His) box polypeptide 21	1.59	NM_004728
calcium channel, voltage-dependent, gamma subunit 2	1.59	NM_006078
sphingosine-1-phosphate lyase 1	1.59	NM_003901
ATP-binding cassette, sub-family A (ABC1), member 12	1.59	AB080207
S164 protein	1.59	BF055107
KIAA1849 protein	1.59	AB055760
solute carrier family 7 (cationic amino acid transporter, y+ system), member 8	1.59	AL385343
testudin 3	1.59	NM_016950
KIAA0530 protein	1.59	AA972711
CD64 antigen (leukocyte antigen)	1.59	AF054817
coproporphyrinogen oxidase (coproporphyrin, harderoporphyria)	1.59	NM_000097
protein tyrosine phosphatase type IVA, member 3	1.59	BC003105
KIAA0256 gene product	1.59	DB7445
glyceroldehyde-3-phosphate dehydrogenase	1.59	AL035604
Phtn	1.59	NM_003662
hypothetical protein FLJ20081	1.59	NM_017658
Consensus includes gb:AW149844 /FEA=EST /DB_XREF=gi:197740 /DB_XREF=est:442h10.x1 /CLONE=IMAGE:2620771 /UG=Hs.187140	1.59	AW149844
G protein-coupled receptor	1.59	BC006381
gb:BC006381.1 /DEF=Homo sapiens, clone MGC:13287. mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13287) /DB_XREF=gi:13623544 /FL=gb:BC006381	1.59	BC006381
WAS protein family, member 3	1.59	AB020707
paat	1.59	NM_016566
solute carrier family 4, sodium bicarbonate cotransporter, member 7	1.58	NM_003615
eukaryotic translation initiation factor 1A	1.58	AL079283
Consensus includes gb:AW971205 /FEA=EST /DB_XREF=gi:161050 /DB_XREF=est:EST363293 /UG=Hs.114280	1.58	AW971205
solute carrier family 19 (thiamine transporter), member 2	1.58	AF153330
RAN binding protein 6	1.58	AI123233
tumor protein 63 KDa with strong homology to p53	1.58	AF116771
interleukin 3 (colony-stimulating factor, multiple)	1.58	NM_000588
cold autoinflammatory syndrome 1	1.58	AK027194
M-phase phosphoprotein 1	1.58	NM_016195
par-3 partitioning defective 3 homolog (C. elegans)	1.58	AF196186
hypothetical protein FLJ10512	1.58	AI655902
tumor protein p53 (Li-Fraumeni syndrome)	1.58	NM_000548
etonal homolog 1 (Drosophila)	1.58	NM_005172
small inducible cytokine subfamily A (Cys-Cys), member 16	1.58	NM_004590

colony stimulating factor 1 (macrophage)	M76453	1.58
Consensus includes gb:AL109667.1 /DEF=Homo sapiens mRNA (full length insert cDNA clone EUROMAGE 381867 /FEA=mRNA /DB_XREF=gi:5689815 /UG=Hs.124154	AL109667	1.58
N-myristoyltransferase 1	A1570834	1.58
Consensus includes gb:AL359578.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547N163 (from clone DKFZp547N163) /FEA=mRNA /DB_XREF=gi:5655637 /UG=Hs.3065	AL359578	1.58
KIAA1693 protein	AI869898	1.58
arachidonate 12-lipoxygenase, 12R type	NM_001139	1.58
neurofilin 2	NM_005076	1.58
CD1E antigen, e polypeptide	AF280546	1.58
butyrophilin, subfamily 3, member A3	NM_030893	1.58
HSPC135 protein	NM_006994	1.58
Consensus includes gb:AL049435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp56880220 (from clone DKFZp56880220) /FEA=mRNA /DB_XREF=gi:4500218 /UG=Hs.171	NM_014170	1.58
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12E47)	AL049435	1.58
KIAA0480 gene product	AI655986	1.58
Consensus includes gb:AF52772 /FEA=EST /DB_XREF=gi:5745082 /DB_XREF=est:wr50d05.x1 /CLONE=IMAGE:2491113 /UG=Hs.300865 Homo sapiens clone BCSynL	AW299294	1.58
phosphodiesterase 1B, calmodulin-dependent	NM_000924	1.58
thymine-DNA glycosylase	BF674842	1.58
growth factor independent 1	NM_005263	1.58
interferon induced transmembrane protein 1 (9-27)	A4749101	1.58
carboxylesterase-related protein	NM_016280	1.58
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O	AA758795	1.58
rho/rec guanine nucleotide exchange factor (GEF) 2	NM_004723	1.58
KIAA0296 gene product	NM_014699	1.58
Consensus includes gb:U82306.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938333 /UG=Hs.123081 Homo	U82306	1.58
hypothetical protein FLJ21777	AK025430	1.58
DNA (cytosine-5-methyltransferase 3 beta	NM_006892	1.58
KIAA0475 gene product	NM_014864	1.58
vacuolar protein sorting 45A (yeast)	NM_007258	1.58
uncharacterized hematopoietic stem/progenitor cells protein MDS029	NM_018464	1.58
hypothetical protein FLJ21803	NM_024762	1.58
Pyruvate dehydrogenase complex, lipoyl-containing component X; E3-binding protein	NM_003477	1.58
guanylate kinase 1	AW182392	1.58
general transcription factor IIC, polypeptide 2 (beta subunit, 110kD)	AW194657	1.58
Consensus includes gb:AF257059 /DEF=Homo sapiens prothymosin alpha (PTMA) gene, complete cds /FEA=CDs /DB_XREF=gi:8037944 /UG=Hs.283947 Homo sapiens	AF257059	1.58
Consensus includes gb:AW134608 /FEA=EST /DB_XREF=gi:6138154 /DB_XREF=est:U1-H-B11-abo-e-04-0-U1.s1 /CLONE=IMAGE:2712654 /UG=Hs.168941 ESTs	AW134608	1.58
complement component (3b/4b) receptor 1, including Knops blood group system	X14362	1.58
cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)	NM_001262	1.58
chromosome 16 open reading frame 44	NM_024731	1.58
Consensus includes gb:AA017093 /FEA=EST /DB_XREF=gi:1479257 /DB_XREF=est:ze37h10.s1 /CLONE=IMAGE:361219 /UG=Hs.271780 ESTs	AA017093	1.58
centaurin, alpha 2	NM_018404	1.58
TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor, 135 kD	AI744029	1.58
secreted protein of unknown function	BF440021	1.58
tenilin AC	NM_005572	1.58
fizzled homolog 1 (Drosophila)	AF072872	1.58
disca, large (Drosophila) homolog-associated protein 1	NM_004746	1.58
neuronal specific transcription factor DAT1	AL050152	1.58
U2 small nuclear ribonucleoprotein auxiliary factor (65kD)	AI810156	1.58
Consensus includes gb:AK025065.1 /DEF=Homo sapiens cDNA: FLJ21412 fls, clone COL03987. /FEA=mRNA /DB_XREF=gi:10437504 /UG=Hs.288066 Homo sapiens cd	AK025065	1.58
myosin IC	BE790157	1.58
VLCS-H1 protein	NM_014031	1.58
flavin containing monooxygenase 2	AL021026	1.58
Interleukin 7 receptor	NM_002185	1.58
Nijmegen breakage syndrome 1 (nibrin)	NM_002485	1.58
aquaporin 8, kidney specific	AL137716	1.58
SPRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal)	NM_000346	1.57
hypothetical protein FLJ21240	NM_024847	1.57
succinate-CoA ligase, ADP-forming, beta subunit	NM_003850	1.57
guanine nucleotide binding protein (G protein), alpha 15 (Gq class)	NM_002068	1.57
high-mobility group protein 2-like 1	AL078310	1.57
hypothetical protein FLJ10848	NM_018167	1.57
carbonic anhydrase VA, mitochondrial	NM_001739	1.57
translocated promoter region (to activated MET oncogene)	BF110993	1.57
hypothetical protein FLJ20473	NM_017838	1.57

hypothetical protein LOC37158	1.57	NM_020433
Parkinson disease (autosomal recessive, juvenile) 2, parkin	1.57	NM_004582
hypothetical protein FLJ23188	1.57	NM_025055
nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)	1.57	U09809
mitochondrial triglyceride transfer protein (large polypeptide, 88kD)	1.57	A0823321
Consensus includes gb:AU158608 /FEA=EST /DB_XREF=gi:11020127 /DB_XREF=est:AU158606 /CLONE=PLACE3000406 /UG=Hs.287638 Homo sapiens cDNA FLJ143 AU158606	1.57	AF091627
tumor protein 63 kDa with strong homology to p53	1.57	AF091627
Consensus includes gb:AP035317.1 /DEF=Homo sapiens clone 23892 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2681080 /UG=Hs.91916 Homo sapiens clone 23892, AF035317	1.57	U1020227
PR domain containing 9	1.57	U13699
casepase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	1.57	NM_017973
hypothetical protein FLJ10034	1.57	NM_000311
prion protein (p27-30) (Creutzfeldt-Jakob disease, Gerstmann-Sträussler-Scheinker syndrome, fatal familial Insomnia)	1.57	NM_002264
keratinocyte alpha 1 (imporin alpha 5)	1.57	NM_002530
neurotrophic tyrosine kinase, receptor, type 3	1.57	X51990
T-cell acute lymphocytic leukemia 1	1.57	AW062341
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	1.57	NM_018260
hypothetical protein FLJ10891	1.57	AU118260
actinin, alpha 1	1.57	M95178
Consensus includes gb:AL54245 /FEA=EST /DB_XREF=gi:12894837 /DB_XREF=est:AL54245 /CLONE=CS001082YA02 (5 prime) /UG=Hs.288869 nuclear receptor subunit AL54245	1.57	NM_012334
myosin X	1.57	NM_018097
hypothetical protein FLJ10460	1.57	AK001618
TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 150 kD	1.57	NM_001918
dihydropyrimidine branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex, maple syrup urine disease)	1.57	NM_013248
NTF2-related export protein 1	1.57	AF039198
hairless homolog (mouse)	1.57	AF047033
solute carrier family 4, sodium bicarbonate cotransporter, member 7	1.57	AF221098
Rai guanine nucleotide exchange factor RaiGPS1A	1.57	NM_001648
kallikrein 3, (prostate specific antigen)	1.57	X61079
T cell receptor alpha variable 8-6	1.57	BE350026
zinc finger protein (clone 647)	1.57	NM_024560
hypothetical protein FLJ21983	1.57	AW276522
solute carrier family 8 (neurotransmitter transporter, creatine), member 8	1.57	AB045929
elko-keto reductase family 1, member C4 (chlorocone reductase, 3-alpha hydroxysteroid dehydrogenase, type 1; dihydriol dehydrogenase 4)	1.57	NM_024967
hypothetical protein FLJ11637	1.57	NM_002777
protease 3 (serine protease, neutrophil, Wegener granulomatosis autoantigen)	1.57	NM_002777
Consensus includes gb:BE88744 /FEA=EST /DB_XREF=gi:10345354 /DB_XREF=est:501513656F1 /CLONE=IMAGE:3915133 /UG=Hs.169274 ESTs. Highly similar to IF BE88744	1.57	AF0708130
Consensus includes gb:AU0708130 /FEA=EST /DB_XREF=est:AU0708130 /CLONE=ADCA05 /UG=Hs.283655 Homo sapiens scavenger receptor AF0708130	1.57	AF0708130
Consensus includes gb:AL0333403 /DEF=Human DNA sequence from clone 80D7 on chromosome Xq25-26.3 Contains F9 (coagulation factor IX (plasma thromboplastin co	1.57	AL033403
KLHL1547 protein	1.57	A067426
KIAA1547 protein	1.57	NM_004685
Consensus includes gb:AW663632 /FEA=EST /DB_XREF=gi:7456210 /DB_XREF=est:h12a03.x1 /CLONE=IMAGE:2981548 /UG=Hs.285825 Homo sapiens mRNA: cDNA AW663632	1.57	AB347136
telomeric repeat binding factor (NIMA-interacting) 1	1.57	AK023321
chromosome 2 open reading frame 6	1.57	A0282485
HLA-B associated transcript 1	1.57	NM_002159
histatin 1	1.57	NM_006683
human epikidymis-specific 3 alpha	1.57	NM_006579
emogamyl binding protein (sterol isomerase)	1.57	U95089
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	1.57	NM_020640
PRP42 homolog	1.57	NM_020640
Consensus includes gb:AL080233.1 /DEF=Homo sapiens mRNA: cDNA DKFZp568L111 (from clone DKFZp568L111) /FEA=mRNA /DB_XREF=gi:5262726 /UG=Hs.32656 /AL080233	1.57	A017382
KIAA1218 protein	1.57	NM_003538
H4 histone family, member A	1.57	NM_003538
Consensus includes gb:BF511231 /FEA=EST /DB_XREF=gi:11594529 /DB_XREF=est:U1-H-B14-aoi-g-11-Q.U1.s1 /CLONE=IMAGE:3085244 /UG=Hs.288582 ESTs. Weakly	1.57	BF511231
spectrin, beta, erythrocytic (includes spherocytosis, clinical type 1)	1.57	NM_000347
splicing factor, arginine/serine-rich 3	1.57	AF107405
uronyl 2-sulfotransferase	1.57	AW418882
vinculin	1.57	NM_014000
platelet-derived growth factor receptor, alpha polypeptide	1.57	M22734
fusion, derived from (t(12;16) malignant liposarcoma	1.57	AW511410
Consensus includes gb:AK021983.1 /DEF=Homo sapiens cDNA FLJ11928 fs, clone HEMBB1000374, /FEA=mRNA /DB_XREF=gi:10433297 /UG=Hs.306617 Homo sapie	1.57	AK021983
enolase 2, (gamma, neuronal)	1.57	NM_001975
Consensus includes gb:BF674749 /FEA=EST /DB_XREF=gi:11948644 /DB_XREF=est:602136893F1 /CLONE=IMAGE:4273270 /UG=Hs.258800 ESTs	1.57	BF674749
hypothetical protein FLJ13197	1.57	NM_024614

157

protein phosphatase 2, regulatory subunit B (B56), alpha isoform
Consensus includes gb:AK024561.1 /DEF=Homo sapiens cDNA: FLJ20908 fls, clone ADSE00417, /FEA=mRNA /DB_XREF=gi:10436870 /UG=Hs.306689 Homo sapiens c AK024561 1.56
hypothetical protein FLJ21302
growth hormone receptor
Consensus includes gb:AJ251844.1 /DEF=Homo sapiens partial mRNA for MOZCBP chimeric transcript type II, /FEA=mRNA /GEN=MOZCBP /DB_XREF AJ251844 1.56
transmembrane protein 8 (five membrane-spanning domains)
Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene, /FEA=mRNA /DB_XREF=gi:704460 /UG=Hs.285688 H.sapiens clathrin light chain a gene X81636 1.56
pregnancy specific beta-1-glycoprotein 1
HBS1-like (S. cerevisiae)
lymphocyte antigen 8
sirtuin silent mating type information regulation 2 homolog 2 (S. cerevisiae)
leukocyte tyrosine kinase
Consensus includes gb:AL355532 /DEF=Human DNA sequence from clone RP11-487F5 on chromosome 6 Contains the 3 part of the GRIK2 (glutamate receptor, ionotropic) AL355532 1.56
Consensus includes gb:AI083505 /FEA=EST /DB_XREF=gi:5810724 /DB_XREF=est:W48h12.x1 /CLONE=IMAGE:2510855 /UG=Hs.137576 ribosomal protein L34 pseudo AI083505 1.56
hypothetical protein DKFZp782166
FK506 binding protein 6 (38kD)
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1
Consensus includes gb:AV692438 /FEA=EST /DB_XREF=gi:10284299 /DB_XREF=est:AV682436 /CLONE=GKXABE08 /UG=Hs.5848 Homo sapiens mRNA; cDNA DKFZ: AV682436 1.55
zinc finger protein 358
5-hydroxytryptamine (serotonin) receptor 2A
Consensus includes gb:AK024158.1 /DEF=Homo sapiens cDNA FLJ14096 fls, clone MAMMA1000752, /FEA=mRNA /DB_XREF=gi:10436469 /UG=Hs.303289 Homo sapiens AK024158 1.55
endothelin 1
general transcription factor IIH, polypeptide 1 (62KD subunit)
collagen, type I, alpha 1
p53-regulated apoptosis-inducing protein 1
son of sevenless (Drosophila) homolog 2
Consensus includes gb:W85912 /FEA=EST /DB_XREF=gi:1398341 /DB_XREF=est:zh52b12.s1 /CLONE=IMAGE:415679 /UG=Hs.323470 Homo sapiens cDNA: FLJ22106 W85912 1.55
wingless-type MMTV integration site family, member 7B
hypothetical protein DKFZp781D1823
nuclear fragile X mental retardation protein interacting protein 1
hypothetical protein FLJ22490
Consensus includes gb:AV717062 /FEA=EST /DB_XREF=gi:10814214 /DB_XREF=est:AV717062 /CLONE=DCBCEG08 /UG=Hs.274553 Homo sapiens cDNA FLJ10906 fl AV717062 1.55
fibroblast growth factor 20
Consensus includes gb:AL138139 /DEF=Human DNA sequence from clone RP4-7812 on chromosome 6 Contains 3 part of the gene for enhancer of filamentation (HEF1), AL138139 1.55
hypothetical protein FLJ12284
Consensus includes gb:AK023843.1 /DEF=Homo sapiens cDNA FLJ13781 fls, clone PLACE4000465, /FEA=mRNA /DB_XREF=gi:10435902 /UG=Hs.252820 Homo sapiens AK023843 1.55
retinoblastoma (X-linked, juvenile) 1
Consensus includes gb:X16454 /DEF=Human gene for carcinoembryonic antigen subdomains A and B /FEA=CDS /DB_XREF=gi:28852 /UG=Hs.248025 Human gene for c X16454 1.55
gb:BC005233.1 /DEF=Homo sapiens, clone MGC:12257, mRNA, complete cds, /FEA=mRNA /PROD=Unknown (protein for MGC:12257) /DB_XREF=gi:13528868 /UG=Hs. BC005233 1.55
myotubularin related protein 1
hypothetical protein FLJ20874
Rho guanine exchange factor (GEF) 12
fibulin 1 (Marfan syndrome)
solute carrier family 20 (phosphate transporter), member 1
Glutamate receptor interacting protein
kinasin family member 1C
zinc finger protein 257
nucleolin
Consensus includes gb:AK024615.1 /DEF=Homo sapiens cDNA: FLJ20962 fls, clone ADSH00804, /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens c AF254888 1.55
Cdc42 effector protein 4; binder of Rho GTPases 4
Ewing sarcoma breakpoint region 1
hypothetical protein DKFZp761N09121
guanine nucleotide binding protein 13, gamma
cathepsin B
immunoglobulin lambda locus
Consensus includes gb:AV652031 /FEA=EST /DB_XREF=gi:3873045 /DB_XREF=est:AV652031 /CLONE=GLCCWA12 /UG=Hs.282344 Homo sapiens cDNA FLJ13387 fl AV652031 1.55
hypothetical protein FLJ10139
chromosome 15 open reading frame 5
butyrophilin, subfamily 3, member A2
UDP glycosyltransferase 1 family, polypeptide A8
chromosome 11 open reading frame 8
p300/CBP-associated factor

a disintegrin and metalloproteinase domain 10
 lefty homolog-like (mouse)
 gb:BC005220.1 /DEF=Homo sapiens, Similar to chaperonin containing TCP1, subunit 8 (theta), clone MGC:12240, mRNA, complete cds. /FEA=mRNA (PROD=Similar to c
 bordin
 PRO1621 protein
 Rho GDP dissociation inhibitor (GDI) gamma
 CGI-143 protein
 CD24 antigen (small cell lung carcinoma cluster 4 antigen)
 Consensus includes gb:AK024958.1 /DEF=Homo sapiens cDNA: FLJ21305 fts, clone COL02124. /FEA=mRNA /DB_XREF=gi:10437382 /UG=Hs.287658 Homo sapiens c
 hypothetical protein MGC11138
 nerve growth factor receptor (TNFR superfamily, member 16)
 complement component 1, r subcomponent
 synaptotagmin 1
 STAT induced STAT inhibitor 3
 adenosine kinase
 poliovirus receptor
 DKFZF64C1040 protein
 potassium inwardly-rectifying channel, subfamily J, member 9
 TRF-proximal protein
 calpain 3
 3'(2'), 5'-bisphosphate nucleotidase 1
 HEF like Protein
 hairy/enhancer-of-split related with YRPW motif 1
 adenovirus 5 E1A binding protein
 DKFZF64C092 protein
 MHC class I polypeptide-related sequence B
 olfactory receptor, family 12, subfamily D, member 2
 RNA binding motif protein 4
 Consensus includes gb:AA443771 /FEA=EST /DB_XREF=gi:2156446 /DB_XREF=est:zw9508.s1 /CLONE=IMAGE:784743 /UG=Hs.270138 ESTs
 immunoglobulin lambda joining 3
 KIAA0410 gene product
 cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)
 stoned B/TfIIA-alpha/beta-like factor
 centromere protein C 1
 Consensus includes gb:AK021495.1 /DEF=Homo sapiens cDNA FLJ11433 fts, clone HEMBA1001121. /FEA=mRNA /DB_XREF=gi:10432690 /UG=Hs.287415 Homo sapie
 poliovirus receptor-related 2 (herpesvirus entry mediator B)
 KIAA0798 gene product
 transcription elongation factor (SII) elongin A2
 UDP-N-acetyl-alpha-D-galactosamine-6-phosphate N-acetylgalactosaminyltransferase 3 (GalNAc-T3)
 Consensus includes gb:AK021514.1 /DEF=Homo sapiens cDNA FLJ11452 fts, clone HEMBA1001435. /FEA=mRNA /DB_XREF=gi:10432710 /UG=Hs.148598 Homo sapie
 synaptonemal complex protein 1
 a disintegrin and metalloproteinase domain 22
 Consensus includes gb:AL049252.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564D183 (from clone DKFZp564D193). /FEA=mRNA /DB_XREF=gi:4498983 /UG=Hs.3020 AL049252
 resident helix loop helix 2
 Consensus includes gb:AL080106.1 /DEF=Homo sapiens mRNA: cDNA DKFZp568O053 (from clone DKEZp568O053). /FEA=mRNA /DB_XREF=gi:5262531 /UG=Hs.3063 AL080106
 Consensus includes gb:AK022120.1 /DEF=Homo sapiens cDNA FLJ12058 fts, clone HEMBB1002092. /FEA=mRNA /DB_XREF=gi:10433445 /UG=Hs.286688 Homo sapie
 integrin-binding sialoprotein (bone sialoprotein, bone sialoprotein II)
 Cdc42 effector protein 3
 tumor protein p73
 lyxyl oxidase-like 1
 gb:1.14458.1 /DEF=Human Ig rearranged kappa-chain gene V-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398492 /FL=gb:L.14458.1
 sulfate carrier family 7 (sulfate amino acid transporter, y+ system), member 8
 glycolipid transfer protein
 CATT tumor necrosis factor conversion 1
 small inducible cytokine A2 (monocyte chemotactic protein 1)
 hypothetical protein FLJ22332
 transcription factor eta
 hypothetical protein MGC2474
 leucine zipper-like protein
 Consensus includes gb:AF038193.1 /DEF=Homo sapiens clone 23608 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2795913 /UG=Hs.6220 Homo sapiens clone 23608 m
 chemokine (C-C motif) receptor-like 2
 reticulocalbin 2, EF-hand calcium binding domain

DKFZp434P211 protein 1.54
 early lymphoid activation protein 1.54
 hypothetical protein PRO2032 1.54
 vitamin A responsive; cytoskeleton related 1.54
 putative purinergic receptor 1.54
 potassium voltage-gated channel, KQT-like subfamily, member 1 1.54
 A kinase (PRKA) anchor protein 5 1.54
 integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12) 1.54
 protein A 1.54
 Consensus includes gb:AL109671.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 29222. /FEA=mRNA /DB_XREF=gi:5689802 /UG=Hs.140833 /AL109671 1.54
 hypothetical protein 1.54
 CSR1 protein 1.54
 meprin A, beta 1.54
 gamma-glutamyltransferase 1 1.54
 integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12) 1.54
 hypothetical protein FLJ10512 1.54
 fibronectin 1 1.54
 apolipoprotein C-II 1.54
 M-phase phosphoprotein 10 (U3 small nuclear ribonucleoprotein) 1.54
 Consensus includes gb:AL136094 /DEF=Human DNA sequence from clone RP3-527B10 on chromosome 6q25.1-25.3 Contains a pseudogene similar to HMG (high mobility group) protein 1.54
 bypiase beta 2 1.54
 glutamate dehydrogenase 1 1.54
 fibronectin leucine rich transmembrane protein 2 1.54
 immunoglobulin kappa constant 1.54
 G protein-coupled receptor 22 1.54
 chemokine (C-C motif) receptor 1 1.54
 bulous pemphigoid antigen 1 (230240KD) 1.54
 hypothetical protein FLJ12439 1.54
 polyadenylation binding protein-interacting protein 1 1.54
 nebulin 1.54
 adrenergic, alpha-1A-, receptor 1.54
 hypothetical protein FLJ20695 1.54
 coxline VI (neuronal) 1.54
 trophoblast glycoprotein 1.54
 immunoglobulin superfamily, member 4 1.54
 activator of CREM in testis 1.54
 Consensus includes gb:AL157428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp781F152 (from clone DKFZp781F152) /FEA=mRNA /DB_XREF=gi:7018458 /UG=Hs.3064 /AL157428 1.54
 Consensus includes gb:AA100250 /FEA=EST /DB_XREF=gi:1647823 /DB_XREF=est:zn4101.s1 /CLONE=IMAGE350003 /UG=Hs.90869 Homo sapiens clones 24622 an AA100250 1.54
 solute carrier family 26, member 4 1.54
 ephrin-A3 1.54
 sodium channel, voltage-gated, type III, alpha polypeptide 1.54
 hypothetical protein FLJ22287 1.54
 KIAA0446 gene product 1.54
 arachidonate 15-lipoxygenase 1.54
 SH3-domain protein 5 (pontin) 1.54
 DKFZP588L151 protein 1.54
 KIAA1091 protein 1.54
 Consensus includes gb:AK022192.1 /DEF=Homo sapiens cDNA FLJ12130 fis, clone MAMMA1000251. /FEA=mRNA /DB_XREF=gi:10433534 /UG=Hs.319567 Homo sapie 1.53
 hypothetical protein FLJ20502 1.53
 zinc finger protein 236 1.53
 synaptopodin 1.53
 sodium channel, voltage-gated, type XII, alpha polypeptide 1.53
 Consensus includes gb:AL117520.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P2435 (from clone DKFZp434P2435) /FEA=mRNA /DB_XREF=gi:5912036 /UG=Hs.304 /AL117520 1.53
 amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2) 1.53
 hypothetical protein DKFZp564K142 similar to implantation-associated protein 1.53
 endothelial PAS domain protein 1 1.53
 insulin-like growth factor binding protein, acid labile subunit 1.53
 POU domain, class 6, transcription factor 1 1.53
 nucleobindin 2 1.53
 solute carrier family 22 (organic cation transporter), member 2 1.53
 nuclear LIM interactor-interacting factor 1.53
 growth differentiation factor 2 1.53
 gb:BC003528.1 /DEF=Homo sapiens, clone MGC:10782, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10782) /DB_XREF=gi:13087626 /UG=Hs. BC003528 1.53

Consensus includes gb:AU147295 /FEA=EST /DB_XREF=gi:11008816 /DB_XREF=est:AU147295 /CLONE=MAMMA1000264 /UG=Hs.288575 Homo sapiens cDNA FLJ14 AU147295
protein phosphatase 1, regulatory (inhibitor) subunit 12A
interferon, alpha-inducible protein 27
A817061 1.53
NM_005532 1.53
NM_006716 1.53
SRY (sex determining region Y)-box 11
Consensus includes gb:U81981 /DEF=Homo sapiens amiloride-sensitive epithelial sodium channel alpha subunit alpha ENAC-1 and alpha ENAC-2 (SCNN1A) genes, allent
U81981 1.53
AW157202 1.53
L22431 1.53
NM_002042 1.53
NM_025109 1.53
NM_022754 1.53
NM_000599 1.53
AL587302 1.53
AW138827 1.53
NM_014618 1.53
Consensus includes gb:AK026493.1 /DEF=Homo sapiens cDNA: FLJ22840 fls, clone KAJA4709 /FEA=mRNA /DB_XREF=gi:10433368 /UG=Hs.287293 Homo sapiens cDI
AK026493 1.53
AA639705 1.53
NM_017636 1.53
AK021881 1.53
AK021842 1.53
NM_016412 1.53
NM_001523 1.53
NM_018357 1.53
AF220130 1.53
BE982456 1.53
AW024085 1.53
AI346350 1.53
X65232 1.53
Consensus includes gb:AK000220.1 /DEF=Homo sapiens cDNA FLJ20213 fls, clone COLF1971, highly similar to AB007921 Homo sapiens mRNA for KIAA0452 protein, fl
AK000220 1.53
AK033377 1.53
Consensus includes gb:A1697803 /FEA=EST /DB_XREF=gi:4985703 /DB_XREF=est:wa17605.x1 /CLONE=IMAGE2341376 /UG=Hs.114963 Homo Sapiens (clone 83B3E
A1697803 1.53
NM_004750 1.53
NM_024821 1.53
NM_000830 1.53
AF107493 1.53
NM_018696 1.53
NM_018058 1.53
NM_018229 1.53
BE045816 1.53
U48752 1.53
NM_000540 1.53
AI824831 1.53
NM_030970 1.53
NM_005071 1.53
AU121975 1.53
NM_016649 1.53
NM_014887 1.53
NM_030577 1.53
Consensus includes gb:AK025180.1 /DEF=Homo sapiens cDNA: FLJ21527 fls, clone COL05961 /FEA=mRNA /DB_XREF=gi:10437643 /UG=Hs.306779 Homo sapiens cC
AK025180 1.53
NM_013368 1.53
NM_003970 1.53
NM_001153 1.53
NM_018042 1.53
AF181985 1.53
Consensus includes gb:AK022006.1 /DEF=Homo sapiens cDNA FLJ11944 fls, clone HEMMB1000684 /FEA=mRNA /DB_XREF=gi:10433318 /UG=Hs.300771 Homo sapie
AK022006 1.53
NM_002693 1.53
NM_001486 1.53
U57059 1.52
U57843 1.52
AF083810 1.52
NM_000905 1.52
AF052095 1.52
M31222 1.52
Consensus includes gb:AF052095.1 /DEF=Homo sapiens clone 23911 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3360401 /UG=Hs.167344 Homo sapiens clone 23911
AF052095 1.52
transcription factor 3 (E2A Immunoglobulin enhancer binding factors E12/E47)
DNA segment, numerous copies, expressed probes (GS1 gene)
NM_012080 1.52

KIAA0974 protein
 Consensus includes gb:AL117546.1 /DEF=Homo sapiens mRNA; cDNA DKFp5586C1923 (from clone DKFp5586C1923). /FEA=mRNA /DB_XREF=gi:5912079 /UG=Hs.24
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1
 killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2
 aggrscan 1 (chondroin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122)
 cutaneous T-cell lymphoma tumor antigen so70-2
 hypothetical protein
 sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)
 thioaurine S-methyltransferase
 Consensus includes gb:AL117451.1 /DEF=Homo sapiens mRNA; cDNA DKFp5586E2317 (from clone DKFp5586E2317). /FEA=mRNA /DB_XREF=gi:5911901 /UG=Hs.28
 hypothetical protein FLJ20716
 hypothetical protein PRO1446
 protocadherin 1 (cadherin-like 1)
 APG3 autophagy 5-like (S. cerevisiae)
 gb:L14453.1 /DEF=Human ig rearranged nu-chain gene V-H-D-N-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398494 /FL=gb:L14453.1
 docking protein 2, 50kD
 cell growth regulatory with EF-hand domain
 LAP (leucine-rich repeats and PDZ) and no PDZ protein
 class-I MHC-restricted T cell associated molecule
 likely homolog of rat kinase D-interacting substance of 220 kDa
 Ig superfamily protein
 phosphodiesterase 1C, calmodulin-dependent (70kD)
 phosphothioesterase-3-kinase, regulatory subunit, polypeptide 2 (p85 beta)
 interferon (alpha, beta and omega) receptor 2
 Consensus includes gb:AA335056 /FEA=EST /DB_XREF=gi:2279319 /DB_XREF=est:nf84a10.51 /CLONE=IMAGE:926566 /UG=Hs.286553 Homo sapiens, clone IMAGE:3
 Consensus includes gb:AL181956.1 /DEF=Homo sapiens mRNA; cDNA DKFp761A17121 (from clone DKFp761A17121). /FEA=mRNA /DB_XREF=gi:7328008 /UG=Hs.286553
 Consensus includes gb:AK001065.1 /DEF=Homo sapiens cDNA FLJ10203 fit, clone HEMBA1004930, moderately similar to 26S PROTEASOME SUBUNIT S5B. /FEA=ntf AK001065
 chromosome X open reading frame 6
 pleckstrin homology, Sec7 and coiled-coil domains, binding protein
 wingless-type MMTV integration site family, member 8B
 KIAA0944 protein
 Consensus includes gb:A927067 /FEA=EST /DB_XREF=gi:5663031 /DB_XREF=est:wo87f01.x1 /CLONE=IMAGE:2462329 /UG=Hs.289068 Homo sapiens cDNA FLJ119
 vinculin
 lysosomal apyrase-like 1
 glycerol-3-phosphate dehydrogenase 1 (soluble)
 SH3-domain, GRB2-like, endophilin B2
 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
 hypothetical protein FLJ14075
 hypothetical protein FLJ20313
 guanine nucleotide binding protein 4
 tetraspan NET-6 protein
 Consensus includes gb:AUI56204 /FEA=EST /DB_XREF=gi:11017725 /DB_XREF=est:AUI56204 /CLONE=PLACE:1002583 /UG=Hs.306651 Homo sapiens cDNA FLJ134
 Consensus includes gb:AV702306 /FEA=EST /DB_XREF=gi:10718636 /DB_XREF=est:AV702306 /CLONE=ADBDID05 /UG=Hs.321211 ESTs, Highly similar to zinc finger 1
 protein tyrosine phosphatase, receptor type, D
 discs, large (Drosophila), homolog-associated protein 1
 retinoic acid induced 17
 pleckstrin homology-like domain, family A, member 3
 Consensus includes gb:D25272.1 /DEF=Homo sapiens mRNA, clone:RES4-16. /FEA=mRNA /DB_XREF=gi:464187 /UG=Hs.174225 Homo sapiens mRNA, clone:RES4-16
 heterogeneous nuclear ribonucleoprotein L
 homolog of mouse BMP-2 inducible kinase
 dynamin 1
 GDNF family receptor alpha 2
 dopamine receptor D1
 gb:BC005043.1 /DEF=Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA,
 multiple inositol polyphosphate histidine phosphatase, 1
 nuclear receptor coactivator 1
 fibroblast growth factor 18
 G protein-coupled receptor 20
 olfactory receptor, family 7, subfamily E, member 2 pseudogene
 malignancy-associated protein
 NADH:ubiquinone oxidoreductase MLRQ subunit homolog
 homeo box (H8 family) 1
 basic-helix-loop-helix-PAS protein

sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida binding)
 Consensus includes gb:AW007988 /FEA=EST /DB_XREF=gi:5856766 /DB_XREF=est:vw48a04.x1 /CLONE=IMAGE:2532750 /UG=Hs.233299 ESTs
 clusterin-like 1 (retinol)
 parathyroid hormone-like hormone
 melanoma antigen, family E, 1, cancer/testis specific
 PL6 protein
 hypothetical protein DKFZp470146
 Consensus includes gb:AF052108.1 /DEF=Homo sapiens clone 23687 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360415 /UG=Hs.12513 Homo sapiens clone 23687 /AF052108
 actinin A receptor, type IB
 chromobox homolog 5 (HP1 alpha homolog, Drosophila)
 neuronal cell adhesion molecule
 Consensus includes gb:AF308291.1 /DEF=Homo sapiens serologically defined breast cancer antigen NY-BR-46 mRNA, partial cds. /FEA=mRNA /PROD=serologically def AF308291
 interleukin receptor matrix proteoglycan 1
 rap guanine nucleotide exchange factor
 keratin 14 (epidermolysis bullosa simplex, Dowling-Meara, Koebner)
 nardin 4
 STAT induced STAT inhibitor 3
 chitinase 3-like 1 (cartilage glycoprotein-39)
 Consensus includes gb:AK026720.1 /DEF=Homo sapiens cDNA: FLJ23067 fls, clone LNG04993. /FEA=mRNA /DB_XREF=gi:10439638 /UG=Hs.117167 Homo sapiens cDNA: AK026720
 KIAA0738 gene product
 a disintegrin and metalloproteinase domain 3a (cyrtestin 1)
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 1
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 10
 protein-O-mannosyltransferase 1
 adrenomedullin beta-1, receptor
 Consensus includes gb:AU159412 /FEA=EST /DB_XREF=gi:11020933 /DB_XREF=est:AU159412 /CLONE=THYRO1001602 /UG=Hs.296756 Homo sapiens cDNA: FLJ141: AU159412
 killer cell lectin-like receptor subfamily C, member 4
 nectin 3
 Consensus includes gb:AF131813.1 /DEF=Homo sapiens clone 24970 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4406647 /UG=Hs.21258 Homo sapiens cDNA: FLJ131: AF131813
 DnaJ (Hsp40) homolog, subfamily C, member 8
 gb:AF116914.1 /DEF=Homo sapiens PRO3078 mRNA, complete cds. /FEA=mRNA /PROD=PRO3078 /DB_XREF=gi:7770264 /UG=Hs.306573 Homo sapiens PRO3078 m AF116914
 Nck, Ash and phospholipase C binding protein
 hypothetical protein FLJ22688
 hypothetical protein FLJ22081
 kinase suppressor of ras
 likely ortholog of mouse early B-cell factor 2
 alcohol dehydrogenase 1B (class I), beta polypeptide
 DKFZP434C091 protein
 nuclear receptor interacting protein 1
 nuclear factor I/B
 pleckstrin homology, Sec7 and coiled/coiled domains 1 (cytohesin 1)
 laminin, gamma 2 (nbsin (100kD), kalinin (105kD), BM600 (100kD), Herlitz junctional epidermolysis bullosa))
 predicted osteoblast protein
 Bacillus subtilis /REF=AF138424 /DEF=B subtilis dapB, [ojF, [ojG genes corresponding to nucleotides 1435-1646 of L38424 /LEN=1931 (-5, -M, -3 represent transcript regions L38424
 peptidyl/prolyl isomerase C (cyclophilin C)
 hypothetical protein FLJ11210
 tumor-associated calcium signal transducer 2
 Immune dysregulation, polyendocrinopathy, enteropathy, X-linked
 cyclin, basic protein of sperm head cytoskeleton 1
 olfactory receptor, family 51, subfamily E, member 2
 defensin, alpha 4, corticostatin
 Consensus includes gb:AU145135 /FEA=EST /DB_XREF=gi:11006656 /DB_XREF=est:AU145135 /CLONE=HEMBA1003989 /UG=Hs.274266 Homo sapiens cDNA: FLJ10: AU145135
 complement component (Bb4b) receptor 1, including Knops blood group system
 DKFZP434M131 protein
 G-protein coupled receptor
 ceruloplasmin (ferroxidase)
 hypothetical protein FLJ11038
 insulin-like growth factor 1 (somatomedin C)
 5-hydroxytryptamine (serotonin) receptor 4
 myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila): translocated to, 4
 Consensus includes gb:BE677757 /FEA=EST /DB_XREF=gi:10039372 /DB_XREF=est:75909.x1 /CLONE=IMAGE:3299008 /UG=Hs.311561 ESTs, Weakly similar to CY, BE677757
 hypothetical protein FLJ22099
 Consensus includes gb:AK025325.1 /DEF=Homo sapiens cDNA: FLJ21672 fls, clone COL09025. /FEA=mRNA /DB_XREF=gi:10437818 /UG=Hs.306801 Homo sapiens cDNA: AK025325

thyroid hormone receptor-associated protein, 240 kDa subunit
 hypothetical protein FLJ22870
 scaffold attachment factor B
 hypothetical protein PRO2176
 zinc finger protein 345
 SH3-domain protein 5 (ponsin)
 PDZ-73 protein
 Smc2-related CBP activator protein
 gb:U79250.1 /DEF=Human glycerol-3-phosphate dehydrogenase mRNA, complete cds. /FEA=mRNA /PROD=glycerol-3-phosphate dehydrogenase /DB_XREF=gi:1806827 /U79250
 KIAA0548 protein
 gb:BC002778.1 /DEF=Homo sapiens. Similar to myosin light chain 2, precursor lymphocyte-specific, clone MGC-3479, mRNA, complete cds. /FEA=mRNA /PROD=Similar
 guanine nucleotide binding protein (G protein), alpha 11 (Gq class)
 Consensus includes gb:AF131777.1 /DEF=Homo sapiens clone 25081 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4408602 /UG=Hs.183475 Homo sapiens clone 25081 AF131777
 A kinase (PRKA) anchor protein 1
 KIAA1105 protein
 ecdonucleoside triphosphate diphosphohydrolase 3
 hypothetical protein FLJ20623
 KIAA0557 protein
 DMC1 dosage suppressor of meck1 homolog, meiosis-specific homologous recombination (yeast)
 LOC89231
 ectodermal dysplasia 1, anhidrotic
 ATP-binding cassette, sub-family B (MDR/TAP), member 11
 ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D)
 Consensus includes gb:AF178300 /FEA=EST /DB_XREF=gi:4371526 /DB_XREF=est:tm39e01.x1 /CLONE=IMAGE:2160504 /UG=Hs.192789 ESTs, Weakly similar to ALU AF178300
 hypothetical protein FLJ10803
 purinergic receptor P2Y, G-protein coupled, 2
 likely ortholog of mouse SRY-box containing gene 17
 medium-chain acyl-CoA synthetase
 KIAA0429 gene product
 arsenate resistance protein ARS2
 hypothetical protein FLJ20700
 ATPase, Class V, type 10C
 PRO3053 protein
 Consensus includes gb:AL080086.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J1762 (from clone DKFZp564J1762). /FEA=mRNA /DB_XREF=gi:5262502 /UG=Hs.274 AL080086
 mitogen-activated protein kinase kinase kinase 12
 protein kinase, cGMP-dependent, type 1
 follistatin-like 1
 Consensus includes gb:AL035087 /DEF=Human DNA sequence from clone 170F5 on chromosome Xq22.3-24. Contains an HMGI (high-mobility group (nonhistone) chromatin) domain
 Consensus includes gb:AK024896.1 /DEF=Homo sapiens cDNA: FLJ21243 (fs, clone COL01164). /FEA=mRNA /DB_XREF=gi:10437310 /UG=Hs.288016 Homo sapiens cDNA: AK024896
 cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)
 hypothetical protein FLJ11583
 solute carrier family 22 (organic cation transporter), member 5
 Consensus includes gb:AL359553 /DEF=Human DNA sequence from clone RPS-871G17 on chromosome 1. Contains ESTs, STSs and GSSs. Contains the HAQZ gene for AL359553
 Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722). /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.33 AL080112
 Consensus includes gb:AK022331.1 /DEF=Homo sapiens cDNA: FLJ12269 (fs, clone MAMMA1001635). /FEA=mRNA /DB_XREF=gi:10433706 /UG=Hs.283925 Homo sapiens cDNA: AK022331
 Consensus includes gb:U80770.1 /DEF=Human EST clone 251800 mariner transposon Hamar1 sequence. /FEA=mRNA /DB_XREF=gi:2921608 /UG=Hs.42182 Human EST: U80770
 KIAA0877 gene product
 ecotropic viral integration site 2B
 low density lipoprotein-related protein 2
 2-5-oligoadenylate synthetase-like
 hypothetical protein FLJ10718
 N-acetylated alpha-linked acidic dipeptidase-like; (LEAL DIPEPTIDYLPEPTIDASE
 aldehyde dehydrogenase 3 family, member B1
 hypothetical protein
 hypothetical protein MGC13053
 Consensus includes gb:AA131826 /FEA=EST /DB_XREF=gi:1693379 /DB_XREF=est:z37f06.s1 /CLONE=IMAGE:504131 /UG=Hs.107164 spectrin, beta, non-erythrocytic
 KIAA0227 protein
 toll-like receptor 2
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 4
 cell division cycle 2-like 2
 CREBBP/EP300 inhibitory protein 1
 RAB2, member RAS oncogene family
 chromosome 3p21.1 gene sequence

145

uroplakin 18	NM_008952	1.50
LAT1-3TM protein	AC003007	1.50
synaptotagmin V	NM_003180	1.50
period homolog 3 (Drosophila)	Z98884	1.50
tumor necrosis factor (ligand) superfamily, member 9	NM_003811	1.50
hypothetical protein FLJ22269	AK025922	1.50
solute carrier family 12 (potassium/chloride transporters), member 4	A017690	1.50
Fzr1 protein	A173260	1.50
VAMP (vesicle-associated membrane protein)-associated protein B and C	NM_004738	1.50
Cluster incl. A1557319:PT2.1, 16, F08.7 Homo sapiens cDNA, 3' end, clone_end=3 /gb=AL1557319/gi=4489682 /ug=Hs.21921 /len=876	A1557319	1.50
Consensus includes gb:AL080068.1 /DEF=Homo sapiens mRNA, cDNA DKFZp564J062 (from clone DKFZp564J062), /FEA=mRNA /DB_XREF=gi:5282475 /UG=Hs.27253	AL080068	1.50
apolipoprotein B (including Ag(x) antigen)	NM_000384	1.50
chimerin (chimerin) 2	L29126	1.50
SEC24 related gene family, member A (S. cerevisiae)	AJ131244	1.50
glycine amidinotransferase (L-arginine:glycine amidinotransferase)	NM_001482	1.50
autism-related protein 1	AK025286	1.50
adenosine monophosphate deaminase 2 (isoform L)	A016249	1.50
hypothetical protein FLJ23231	NM_025079	1.50
Consensus includes gb:AL137292.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434L1016 (from clone DKFZp434L1016), /FEA=mRNA /DB_XREF=gi:6807752 /UG=Hs.274	AL137292	1.50
Consensus includes gb:AK022322.1 /DEF=Homo sapiens cDNA FLJ12260 fs. clone MAMMA100155.1, /FEA=mRNA /DB_XREF=gi:10433693 /UG=Hs.306634 Homo sapie	AK022322	1.50
siaphorin (gpl.115, leukostallin, CD43)	X69502	1.50
gaslin-releasing peptide	NM_002091	1.50
hypothetical protein DKFZp547H025	NM_020181	1.50
zinc finger protein 222	NM_013360	1.50
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 5	J07639	1.50
RYK receptor-like tyrosine kinase	NM_002956	1.50
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	U55209	1.50
PRP4/STKWD splicing factor	A118402	1.50
prothymosin, alpha (gene sequence 28)	BF686442	1.50
hypothetical protein FLJ23047	NM_024548	1.50
sulfotransferase family, cytosolic, 2B, member 1	NM_004605	1.50
diaphanous homolog 2 (Drosophila)	NM_007309	1.50
Consensus includes gb:BE892698 /FEA=EST /DB_XREF=gi:10353302 /DB_XREF=est6014339 /TF1 /CLONE=IMAGE:3918989 /UG=Hs.129709 Homo sapiens chromoso	BE892698	1.50
Consensus includes gb:N30878 /FEA=EST /DB_XREF=gi:1149398 /DB_XREF=est:YW7605.s1 /CLONE=IMAGE:238129 /UG=Hs.82276 trichohyalin	N30878	1.50
filipin 1	AA507012	1.50
platelet-derived growth factor receptor, beta polypeptide	NM_002609	1.50
coagulation factor II (thrombin)	NM_000506	1.50
ankyrin 3, node of Ranvier (ankyrin G)	NM_001149	1.50
tetraspan 5	AA748177	1.50
Consensus includes gb:A1825798 /FEA=EST /DB_XREF=gi:54464689 /DB_XREF=est:td18e08.x1 /CLONE=IMAGE:2076038 /UG=Hs.211388 Homo sapiens BAC clone CTE	A1825798	1.50
chromosome 11 open reading frame 9	AC004770	1.50
solute carrier family 22 (organic anion transporter), member 7	AF210455	1.50
pp21 homolog	NM_016303	1.50
hypothetical protein FLJ10261	NM_018043	1.50
oligodactyl receptor, family 3, subfamily A, member 2	NM_002551	1.49
FXD domain-containing ion transport regulator 2	NM_001680	1.49
TSPY-like	AL136629	1.49
Consensus includes gb:AL359563.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434D0326 (from clone DKFZp434D0326), /FEA=mRNA /DB_XREF=gi:8655619 /UG=Hs.30	AL359563	1.49
PTD011 protein	NM_014051	1.49
Consensus includes gb:AL137606.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434H0211 (from clone DKFZp434H0211), /FEA=mRNA /DB_XREF=gi:6808354 /UG=Hs.15	AL137606	1.49
flap structure-specific endonuclease 1	BC000323	1.49
glucagon-like peptide 2 receptor	NM_004246	1.49
old astrocyte specifically induced substance	BC328407	1.49
promyelocytic leukemia	AF230410	1.49
glutamate-ammmonia ligase (glutamine synthase)	U08626	1.49
H3 histone family, member K	NM_003538	1.49
KIA00881 protein	AB023198	1.49
erythrocyte membrane protein band 4.1-like 1	AA573523	1.49
major histocompatibility complex, class II, DQ beta 1	M24364	1.49
gb:AF-098641.1 /DEF=Homo sapiens CD44 isoform RC (CD44) mRNA, complete cds. /FEA=mRNA /GEN=CD44 /PROD=CD44 isoform RC /DB_XREF=gi:3832517 /UG=Hs.1	AF098641	1.49
siRNA silencing type information regulation 2 homolog 4 (S. cerevisiae)	NM_012240	1.49
Consensus includes gb:U05589.1 /DEF=Human ribosomal protein S1 homolog mRNA, partial cds. /FEA=mRNA /DB_XREF=gi:497001 /UG=Hs.371 ribosomal protein S1-III	U05589	1.49
myristoylated alanine-rich protein kinase C substrate	AA770596	1.49

calcitonin receptor
desmoplakin (DPI, DPII)
hypothetical protein MGC3178
thymase 1, mast cell
protein phosphatase 1, regulatory (inhibitor) subunit 1A
fibroblast activation protein, alpha
Consensus includes gb:AK000122.1 /DEF=Homo sapiens cDNA FLJ20115 fs, clone COL05594, /FEA=mRNA /DB_XREF=gi:7020002 /UG=Hs.306382 Homo sapiens cDNA AK000122
T-box 2
hypothetical protein FLJ1280
Consensus includes gb:BF511164 /FEA=EST /DB_XREF=gi:11594462 /DB_XREF=est:U1-H-B14-aoi-a-09-O-U1.s1 /CLONE=IMAGE:3084952 /UG=Hs.293309 ESTs
transmembrane 6 superfamily member 2
KIA0953 protein
macrophage stimulating 1 receptor (c-met-related tyrosine kinase)
erythrocyte protein band 4.1-like 4
Consensus includes gb:X63749 /DEF=H sapiens DNA for Rod transducin alpha-subunit /FEA=mRNA /DB_XREF=gi:37049 /UG=Hs.51147 guanine nucleotide binding prote
S100 calcium binding protein A8 (calgranulin A)
hypothetical protein FLJ20043
protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)
keratin 7
mitochondrial ribosomal protein L9
paired mesoderm homeo box 1
hypothetical protein MGC3136
ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)
TATA box binding protein (TBP)-associated factor, RNA polymerase I, B, 63KD
NK6 transcription factor homolog A (Drosophila)
Consensus includes gb:A824078 /FEA=EST /DB_XREF=gi:5444749 /DB_XREF=est:w35h06.x1 /CLONE=IMAGE:2404859 /UG=Hs.295448 ESTs, Moderately similar to B
H3 histone family, member J
milk fat globule-EGF factor 8 protein
peroxisomal biogenesis factor 11A
ectodermal microtubule-associated protein-like
adrenergic, alpha-1A, receptor
peptidylglycine alpha-amidating monooxygenase
centaurin, delta 1
caldesmon 1
Consensus includes gb:AL353942.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L23121 (from clone DKFZp761L23121) /FEA=mRNA /DB_XREF=gi:7669881 /UG=Hs.3
hypothetical protein FLJ11856
sulfolipase family, cytosolic, 1C, member 1
phosphate cytidylyltransferase 1, choline, beta isoform
immunoglobulin kappa constant
POU domain, class 4, transcription factor 3
methyl CpG binding protein 2 (Rett syndrome)
prostate androgen-regulated transcript 1
cytochrome P450, subfamily IIC (methoxytoxin 4-hydroxylase), polypeptide 9
interleukin 10 receptor, alpha
KIA0553 protein
Consensus includes gb:A054381 /FEA=EST /DB_XREF=gi:3322168 /DB_XREF=est:q8409.x1 /CLONE=IMAGE:1861265 /UG=Hs.176139 ESTs
Consensus includes gb:AA830143 /FEA=EST /DB_XREF=gi:2903242 /DB_XREF=est:oc44e11.s1 /CLONE=IMAGE:1352588 /UG=Hs.15535 Homo sapiens clone 24582 m
COP9 homolog
Consensus includes gb:AA830143 /FEA=EST /DB_XREF=gi:2903242 /DB_XREF=est:oc44e11.s1 /CLONE=IMAGE:1352588 /UG=Hs.15535 Homo sapiens clone 24582 m
calponin 1, basic, smooth muscle
Consensus includes gb:AU147402 /FEA=EST /DB_XREF=gi:11008923 /DB_XREF=est:AU147402 /CLONE=NAMMA1000567 /UG=Hs.287496 Homo sapiens cDNA FLJ12
extracellular glycoprotein EMLIN-2 precursor
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1
matrix metalloproteinase 7 (matrilysin, uroline)
LIS1-interacting protein NUDE1, rat homolog
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 6 (RNA helicase, 54KD)
Cp1-interacting zinc finger protein
hypothetical protein FLJ12748
ORF
hypothetical protein MGC12488
signal transducer and activator of transcription 2, 113KD
Consensus includes gb:AL050391.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A181 (from clone DKFZp586A181); partial cds. /FEA=mRNA /GEN=DKFZp586A181 /PI AL050391
gb:UA8722.1 /DEF=Human epidermal growth factor receptor precursor (EGFR) mRNA, complete cds. /FEA=mRNA /PROD=epidermal growth factor receptor precursor /DE UA8722

Consensus includes gb:AL034450 /DEF=Human DNA sequence from clone 11SK14 on chromosome Xq22.3-23 Contains high mobility group protein 2a, ESTs, STS /FEA= AL034450 1.49

Mitochondrial Acyl-CoA Thioesterase 1.49

G protein-coupled receptor kinase 6 AF040751 1.49

Consensus includes gb:AK022235.1 /DEF=Homo sapiens cDNA FLJ12173 fs, clone MAMMA1000696, /FEA=mRNA /DB_XREF=gi:10433588 /UG=Hs.306630 Homo sapiens AK022235 1.49

chondroitin sulfate proteoglycan 2 (versican) NM_004385 1.49

angiotensin 2 NM_001147 1.49

Consensus includes gb:BE85194 /FEA=EST /DB_XREF=est:7g19c05.x1 /CLONE=IMAGE:3306920 /UG=Hs.323053 Homo sapiens mRNA full le BE85194 1.49

myosin VIIA (Usher syndrome 1B (autosomal recessive, severe)) U55208 1.49

thymosin, beta, identified in neuroblastoma cells BF677486 1.49

DnaJ (Hsp40) homolog, subfamily B, member 5 NM_012266 1.49

small inducible cytokine subfamily C, member 2 NM_003175 1.49

hypothetical protein MGC5338 NM_024052 1.49

FE65-LIKE 2 NM_006051 1.49

hypothetical protein FLJ10707 BF590987 1.49

fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson M80634, 1.49

histamine receptor H1 NM_000861 1.48

cylindromatosis (turban tumor syndrome) AW45099 1.48

membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2) NM_024322 1.48

hypothetical protein MGC11266 NM_017618 1.48

hypothetical protein FLJ20006 NM_016201 1.48

Leman coiled-coil protein NM_004381 1.48

cAMP responsive element binding protein-like 1 NM_006315 1.48

ring finger protein 3 H08993 1.48

G2 protein NM_003773 1.48

hyaluronoglucosaminidase 2 NM_001764 1.48

CD18 antigen, b polypeptide NM_022903 1.48

hypothetical protein FLJ12800 NM_021123 1.48

G antigen 7 NM_022352 1.48

epithelial V-like antigen 1 AF275945 1.48

capsase recruitment domain family, member 9 NM_021969 1.48

nuclear receptor subfamily 0, group B, member 2 NM_012333 1.48

c-myc binding protein NM_017572 1.48

G protein-coupled receptor kinase 7 NM_003824 1.48

Fas (TNFRSF6)-associated via death domain BC031169 1.48

calpain 3, (p94) NM_003358 1.48

Consensus includes gb:AL033538 /DEF=Human DNA sequence from clone RP3-477H23 on chromosome 22q12.1-12.2 Contains parts of one or two novel genes, ESTs, S AL033538 1.48

Consensus includes gb:A1911273 /FEA=EST /DB_XREF=est:wg3309.x1 /CLONE=IMAGE:2368921 /UG=Hs.122158 ESTs A1911273 1.48

lipoma HMGC fusion partner NM_005780 1.48

hypothetical protein FLJ21276 NM_024633 1.48

proteodherm alpha 12 AF152308 1.48

Consensus includes gb:AK021457.1 /DEF=Homo sapiens cDNA FLJ11395 fs, clone HEMBA1000594, /FEA=mRNA /DB_XREF=gi:10432649 /UG=Hs.306595 Homo sapiens AK021457 1.48

ubiquitin-like 1 (sentrin) BC006462 1.48

apoptotic protease activating factor NM_013229 1.48

kalikrein 13 AL050220 1.48

CGI-82 protein NM_016010 1.48

protein tyrosine phosphatase, non-receptor type substrate 1 BC004298 1.48

erythrocyte membrane protein band 4.1-like 2 AL049634 1.48

run-related transcription factor 2 NM_001431 1.48

zinc finger protein 144 (Met-18) L40992 1.48

DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 24 BC004858 1.48

clone FLB1727 BG421209 1.48

E3 ubiquitin ligase SMURF1 NM_016414 1.48

potassium inwardly-rectifying channel, subfamily J, member 5 AB046845 1.48

surfactant, pulmonary-associated protein B NM_000890 1.48

chromosome 17 open reading frame 1A J02761 1.48

proliferation-associated ZC4, 38kD NM_006382 1.48

steroid-5-alpha-reductase, alpha polypeptide 2 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 2) BES01952 1.48

epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian) NM_000348 1.48

apolipoprotein L, 2 AW157070 1.48

taused-like kinase 1 AF162666 1.48

potassium voltage-gated channel, Shal-related subfamily, member 2 NM_012281 1.48

Consensus includes gb:BF592058 /FEA=EST /DB_XREF=gi:11684382 /DB_XREF=est:n98g09.x1 /CLONE=IMAGE:3572920 /UG=Hs.86405 Homo sapiens mRNA: cDNA BF592058 1.48

fascin (Strongylocentrotus purpuratus) homolog 3 (actin-bundling protein, testicular)	NM_020369	1.48
glial cell derived neurotrophic factor	NM_000514	1.48
glutathione peroxidase 3 (plasma)	AW149846	1.48
MAQS box transcription enhancer factor 2, polypeptide D (myocyte enhancer factor 2D)	NM_005920	1.48
laminin, alpha 4	U77706	1.48
hypothetical protein MGC3101	AK022844	1.48
ubiquitin specific protease 6 (Tre-2 oncogene)	NM_004505	1.48
Consensus includes gb:A116084 /FEA=EST /DB_XREF=gi:3736702 /DB_XREF=est:q650h05.x1 /CLONE=IMAGE:1742457 /UG=Hs.170212 ESTs, Highly similar to b-TRC A116084	U46006	1.48
cysteine and glycine-rich protein 2	AF047485	1.48
amine oxidase, copper containing 3 (vascular adhesion protein 1)	NM_015961	1.48
ubiquitin associated and SH3 domain containing, A	AF091078	1.48
putative dimethyladenosine transferase	U52428	1.48
Consensus includes gb:U52428 /DEF=Human fatty acid synthase gene, partial cds /FEA=CDS_2 /DB_XREF=gi:1374810 /UG=Hs.83190 fatty acid synthase	AJ240085	1.48
T-cell receptor interacting molecule	LJ20469	1.48
dopamine receptor D3	AI633689	1.48
protein kinase C-like 2	NM_016227	1.48
chromosome 1 open reading frame 9	NM_001720	1.48
bone morphogenetic protein 8 (osteogenic protein 2)	U36189	1.48
Consensus includes gb:B166134 /FEA=EST /DB_XREF=gi:12335349 /DB_XREF=est:602289121F1 /CLONE=IMAGE:4357349 /UG=Hs.250723 FK506 binding protein 12 BF968134	NM_000094	1.48
collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive)	NM_001717	1.48
basonuclin	NM_022777	1.48
hypothetical protein FLJ14117	K00627	1.48
Consensus includes gb:K00627.1 /DEF=human knrl repeat mma (cdna clone pcd-kpnt-8), 3 end. /FEA=mRNA /DB_XREF=gi:337653 /UG=Hs.203776 Human knrl repeat r K00627	M58664	1.48
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	U36189	1.48
P311 protein	U36189	1.48
Consensus includes gb:AB000464.1 /DEF=Homo sapiens mRNA, exon 1.2.3.4, clone:RES4-24A. /FEA=mRNA /DB_XREF=gi:1843395 /UG=Hs.104258 Homo sapiens m AB000464	U11287	1.48
Consensus includes gb:AK026820.1 /DEF=Homo sapiens cDNA: FLJ23187 fls, clone LING09902. /FEA=mRNA /DB_XREF=gi:10439765 /UG=Hs.287730 Homo sapiens cC AK026820	NM_005204	1.48
glutamate receptor, ionotropic, N-methyl D-aspartate 2B	NM_005204	1.48
mitogen-activated protein kinase kinase 8	AL133033	1.48
KIAA1025 protein	AF130057	1.48
gb:AF130057.1 /DEF=Homo sapiens clone FLB5539 PRO1454 mRNA, complete cds. /FEA=mRNA /PROD=PRO1454 /DB_XREF=gi:11493420 /UG=Hs.323288 Homo sap AF130057	U36189	1.48
Consensus includes gb:A1143919 /FEA=EST /DB_XREF=gi:3665728 /DB_XREF=est:q6912.x1 /CLONE=IMAGE:737647 /UG=Hs.193783 Human DNA sequence from c A1143919	U36189	1.48
Consensus includes gb:AK001127.1 /DEF=Homo sapiens cDNA FLJ10265 fls, clone HEMBB1001014. /FEA=mRNA /DB_XREF=gi:7022192 /UG=Hs.253369 Homo sapiens AK001127	U36189	1.48
Consensus includes gb:A1480014 /FEA=EST /DB_XREF=gi:4373182 /DB_XREF=est:tm71b09.x1 /CLONE=IMAGE:2163945 /UG=Hs.13415 Homo sapiens clone 24571 mf A480014	AK024029	1.48
modulator of apoptosis 1	NM_012373	1.48
Consensus includes gb:N21364 /FEA=EST /DB_XREF=gi:1126534 /DB_XREF=est:y50h10.s1 /CLONE=IMAGE:265219 /UG=Hs.26968 Homo sapiens mRNA from chrone N21364	Y15718	1.48
offactory receptor, family 3, subfamily A, member 3	NM_004202	1.48
dystrobrevin, beta	NM_013315	1.48
thymosin, beta 4, Y chromosome	NM_000759	1.48
transmembrane phosphatase with tensin homology	NM_005883	1.48
colony stimulating factor 3 (granulocyte)	NM_001349	1.48
Consensus includes gb:AW188214 /FEA=EST /DB_XREF=gi:6462650 /DB_XREF=est:xj93g09.x1 /CLONE=IMAGE:2664832 /UG=Hs.250389 ESTs, Weakly similar to AS8: AW188214	AF136970	1.48
adenomatous polyposis coli like	AI82559	1.48
aspartyl-IRNA synthetase	NM_014870	1.48
pipecolic acid oxidase	BE148534	1.48
hypothetical protein FLJ11286	U82306	1.48
KIAA475 gene product	U82306	1.48
Consensus includes gb:U82306.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938333 /UG=Hs.123081 Homo sapiens U82306	BE148534	1.48
UV-B repressed sequence, HUR 7	AW156899	1.48
Consensus includes gb:AW156899 /FEA=EST /DB_XREF=gi:6228300 /DB_XREF=est:tau88c11.x1 /CLONE=IMAGE:2783444 /UG=Hs.12520 Homo sapiens clone 23568, ; AW156899	NM_020903	1.48
ubiquitin-specific processing protease	AF76652	1.48
ubiquitin-specific processing protease	NM_003532	1.48
bromodomain adjacent to zinc finger domain, 2A	AF766728	1.47
H3 histone family, member D	BC004546	1.47
Consensus includes gb:AV756729 /FEA=EST /DB_XREF=gi:10914577 /DB_XREF=est:AV756729 /CLONE=BMCACB11 /UG=Hs.317661 ESTs	BG105204	1.47
disrupter of silencing 10	AU149305	1.47
CCR4-NOT transcription complex, subunit 2	AU18736	1.47
matrix metalloproteinase 14 (membrane-inserted)	AB043821	1.47
similar to S. cerevisiae RER1	AA322818	1.47
glycoprotein VI (platelet)	AK022303	1.47
ribosomal protein L17	NM_003151	1.47
Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA FLJ12241 fls, clone MAMMA1001274. /FEA=mRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens signal transducer and activator of transcription 4	NM_003284	1.47
transition protein 1 (during histone to protamine replacement)		

variable charge protein on X with two repeats	NM_016378	1.47
syntrophin	NM_014723	1.47
homologous to yeast nitrogen permease (candidate tumor suppressor)	BC002791	1.47
HCF-binding transcription factor Zhangfei	AA521272	1.47
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9	BC002538	1.47
hypothetical protein FLJ22825	NM_024715	1.47
Consensus includes gb:AK021983.1 /DEF=Homo sapiens cDNA FLJ11921 fls, clone HEMBB1000318, /FEA=mRNA /DB_XREF=gi:10433282 /UG=Hs.306616 Homo sapie	NM_000807	1.47
gamma-aminobutyric acid (GABA) A receptor, alpha 2	NM_005222	1.47
distal-less homeo box 6	NM_024705	1.47
hypothetical protein FLJ13839	Z95118	1.47
zinc finger protein 259, pseudogene	A1742789	1.47
eukaryotic translation initiation factor 4E	AW592563	1.47
KIAA0435 gene product	NM_013445	1.47
glutamate decarboxylase 1 (brain, 67kD)	NM_030076	1.47
tumor stroma and activated macrophage protein DLM-1	NM_030776	1.47
phosphatidylinositol-4-phosphate 5-kinase, type I, alpha	U78576	1.47
KIAA1076 protein	AB028998	1.47
hypothetical protein FLJ23049	NM_024687	1.47
Consensus includes gb:AW972765 /FEA=EST /DB_XREF=gi:8162811 /DB_XREF=estEST384659 /UG=Hs.283299 ESTs	AW972765	1.47
magic roundabout	NM_019055	1.47
chromosome 21 open reading frame 2	NM_004928	1.47
piccolo (presynaptic cytomatrix protein)	AC004082	1.47
KIAA1264 protein	AB040812	1.47
sodium channel, nonvoltage-gated 1, beta (Liddle syndrome)	NM_000336	1.47
cartilage paired-class homeoprotein 1	NM_006982	1.47
DKF ZP56411922 protein	AF245505	1.47
Consensus includes gb:AK001442.1 /DEF=Homo sapiens cDNA FLJ10580 fls, clone NT2RP2003533, /FEA=mRNA /DB_XREF=gi:7022701 /UG=Hs.140402 Homo sapiens	NM_000078	1.47
cholesteryl ester transfer protein, plasma	AA779495	1.47
vacuolar protein sorting 18 (yeast)	AB028156	1.47
CGI-142	NM_008115	1.47
preferentially expressed antigen in melanoma	NM_000115	1.47
endothelin receptor type B	AA527340	1.47
homeo box B8	NM_022561	1.47
Consensus includes gb:AK026439.1 /DEF=Homo sapiens cDNA: FLJ22786 fls, clone KIAA2150, /FEA=mRNA /DB_XREF=gi:10439304 /UG=Hs.306848 Homo sapiens cD	NM_021913	1.47
growth hormone 1	AU14167	1.47
AXL receptor tyrosine kinase	BF218804	1.47
collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant)	NM_006129	1.47
hypothetical protein FLJ20986	AL524035	1.47
bone morphogenetic protein 1	NM_016064	1.47
cell division cycle 2, G1 to S and G2 to M	NM_004019	1.47
gb:NM_016064.1 /DEF=Homo sapiens CGI-131 protein (LOC51648) /PROD=CGI-131 protein /DB_XREF=gi:7706344 /UG=Hs.109	U71268	1.47
dyshoplin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS269, DXS270, DXS272	AV686235	1.47
CCRA-NOT transcription complex, subunit 4	AA827871	1.47
mammian-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor)	NM_021088	1.47
putative homeodomain transcription factor 1	NM_020435	1.47
interferon, alpha 4	AF035302	1.47
connexin46.6	NM_021192	1.47
catenin (cadherin-associated protein), delta 2 (neural plakophilin-related arm-repeat protein)	AV657878	1.47
homeo box D11	NM_018546	1.47
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9	U62467	1.47
hypothetical protein PRO2938	L11669	1.47
tubby homolog (mouse)	AF086641	1.47
tetracycline transporter-like protein	AA211481	1.47
tenascin XB	D38122	1.47
LIM domain binding 3	NM_022360	1.47
tumor necrosis factor (ligand) superfamily, member 6	NM_021837	1.47
human epididymis-specific 3 beta	NM_030568	1.47
hypothetical protein FLJ14084	NM_014698	1.47
hypothetical protein MGC10818	NM_024969	1.47
KIAA0782 gene product	AV704962	1.47
hypothetical protein FLJ11703		
sterol-C4-methyl oxidase-like		

chromosome 4 open reading frame 6	NM_005750	1.47
PCTAIRE protein kinase 1	BE787860	1.47
down-regulator of transcription 1, TBP-binding (negative cofactor 2)	AW516932	1.47
hypothetical protein FLJ20307	NM_017757	1.47
phosphotriesterase related	NM_030664	1.47
interleukin 13 receptor, alpha 2	NM_000640	1.47
microtubule-associated protein tau	J03778	1.47
twist homolog (acrocephalosyndactyl 3; Saethre-Chotzen syndrome) (Drosophila)	X89268	1.47
U2 small nuclear ribonucleoprotein auxiliary factor (65KD)	NM_007279	1.47
AD-O12 protein	NM_016449	1.47
coagulation factor VII (serum prothrombin conversion accelerator)	NM_000131	1.47
insulinoma-associated 1	NM_002196	1.47
masleimind-like 1 (Drosophila)	NM_014757	1.47
a disintegrin and metalloproteinase domain 3a (cyrilysin 1)	X89657	1.47
hypothetical protein FLJ11088	NM_018318	1.47
Consensus includes gb:AK026045.1 /DEF=Homo sapiens cDNA: FLJ22392 fls, clone HRC07868. /FEA=mRNA /DB_XREF=gi:10438761 /UG=Hs.275464 Homo sapiens cDNA: AK026045	AF112372	1.47
TJ6 protein	NM_018519	1.47
Consensus includes gb:AL157398 /DEF=Human DNA sequence from clone RP11-56H7 on chromosome 10. Contains ESTs, STSs and GSSs. Contains the gene for the ne	AL157398	1.47
hypothetical protein PRO2286	AK026097	1.47
hypothetical protein FLJ20013	AI752257	1.47
zinc finger protein 38 (KOX 25)	AU145019	1.47
KIAA1013 protein	U86145	1.47
CD6 antigen	NM_012404	1.47
pp32 related 2	AL047879	1.47
Consensus includes gb:AL047879 /FEA=EST /DB_XREF=gi:4728067 /DB_XREF=est:DKFZp586N1222. s1 /CLONE=IMAGE:942764 /UG=Hs.194251 ESTs, Weakly simi	NM_018327	1.47
hypothetical protein FLJ11112	AL568374	1.47
sepin 6	NM_014787	1.47
DnaJ (Hsp40) homolog, subfamily C, member 6	BC001282	1.47
high-mobility group (nucleosome chromosomal) protein 17-like 3	NM_016341	1.47
pancreas-enriched phospholipase C	NM_021047	1.47
zinc finger protein 253	NM_005197	1.47
checkpoint suppressor 1	L16895	1.47
lysyl oxidase	NM_004678	1.47
variable charge, Y chromosome, 2	AV024467	1.47
DnaJ (Hsp40) homolog, subfamily C, member 4	NM_006725	1.47
CD6 antigen	NM_003052	1.47
solute carrier family 34 (sodium phosphate), member 1	NM_014477	1.47
cdg1 protein	BE677131	1.46
enlyrin repeat domain 6	NM_006494	1.46
Ets2 repressor factor	Y09880	1.46
homeo box D3	AA493300	1.46
Consensus includes gb:AA493300 /FEA=EST /DB_XREF=gi:2223141 /DB_XREF=est:ng97e11.s1 /CLONE=IMAGE:942764 /UG=Hs.104258 Homo sapiens mRNA, exon 1	NM_001468	1.46
G antigen 1	NM_012066	1.46
hypothetical protein	NM_004835	1.46
angiotensin receptor 1	NM_003781	1.46
membrane-bound transcription factor protease, site 1	AF002256	1.46
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4	BC000629	1.46
asparry-IRNA synthetase	NM_025031	1.46
hypothetical protein FLJ21075	AV089415	1.46
secreted frizzled-related protein 4	AF348514	1.46
prothymosin, alpha (gene sequence 28)	NM_001230	1.46
caspase 10, apoptosis-related cysteine protease	AF064105	1.46
CDC14 cell division cycle 14 homolog B (S. cerevisiae)	AA081084	1.46
transcriptional co-activator with PDZ-binding motif (TAZ)	NM_006502	1.46
Consensus includes gb:BC005092.1 /DEF=Homo sapiens, clone IMAGE:3843012, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3843012) /DB_Xi	NM_006544	1.46
SEC10-like 1 (S. cerevisiae)	M89051	1.46
Consensus includes gb:AW204383 /FEA=EST /DB_XREF=gi:6503855 /DB_XREF=est:U1-H-B1-adv-c-03-0-U1.s1 /CLONE=IMAGE:2717957 /UG=Hs.312723 ESTs, Highly	AW204383	1.46
glucokinase (hexokinase 4, maturity onset diabetes of the young 2)	AL098700	1.46
Consensus includes gb:AL098700 /DEF=Human DNA sequence from clone RP3-498B10 on chromosome Xp22.31-22.13. Contains the 3 end of the gene for serine/threoni	NM_018000	1.46
hypothetical protein FLJ10116	NM_020394	1.46
zinc finger protein SBZF3	NM_004951	1.46
Epstein-Barr virus induced gene 2 (lymphocyte-specific G protein-coupled receptor)		
Consensus includes gb:AK000786.1 /DEF=Homo sapiens cDNA FLJ20779 fls, clone COL05077. /FEA=mRNA /DB_XREF=gi:7021088 /UG=Hs.321583 Homo sapiens cDNA AK000786		1.46

Consensus includes gb:AL080130.1 /DEF=Homo sapiens mRNA; cDNA DKFZp34E033 (from clone DKFZp34E033). /FEA=mRNA /DB_XREF=gi:5262569 /UG=Hs.1574i /AL080130 1.46
RNA binding motif, single stranded interacting protein 2 AAS36000 1.46
KIAA1473 protein AB040906 1.46
gb:BC001217.1 /DEF=Homo sapiens. Similar to KIAA0029 protein, clone MSG:767, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0029 protein /DB_XREF=g 1.46
AL136883 1.46
hypothetical protein DKFZp564F013 NM_007173 1.46
protease, serine, 23 NM_014035 1.46
SBB31 protein AF041261 1.46
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 4 NM_003441 1.46
zinc finger protein 141 (clone pHZ-44) NM_014962 1.46
KIAA0952 protein AA365670 1.46
Consensus includes gb:AA365670 /FEA=EST /DB_XREF=gi:2018010 /DB_XREF=est:EST76509 /UG=Hs.261526 ESTs 1.46
Consensus includes gb:AA444520 /FEA=EST /DB_XREF=gi:6986282 /DB_XREF=est:UJH-BIG-akb-h-11-UJH.s /CLONE=IMAGE:2733933 /UG=Hs.256986 ESTs, Moder. AA444520 1.46
vesicle trafficking, beach and anchor containing A1659561 1.46
KIAA0731 protein BC001480 1.46
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 NM_022873 1.46
interferon, alpha-inducible protein (clone IFI-6-16) U82277 1.46
CD414 NM_016570 1.46
midline 2 NM_017447 1.46
Consensus includes gb:AL117447.1 /DEF=Homo sapiens mRNA; cDNA DKFZp568A0617 (from clone DKFZp568A0617). /FEA=mRNA /DB_XREF=gi:5911895 /UG=Hs.30i /AL117447 1.46
secreted frizzled-related protein 1 AF017987 1.46
protein kinase, CAMP-dependent, catalytic, alpha NM_002730 1.46
DKFZP568M1019 protein NM_015603 1.46
angiotensinogen (serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 8) NM_000029 1.46
3-phosphoinositide dependent protein kinase-1 NM_002613 1.46
Consensus includes gb:AK021474.1 /DEF=Homo sapiens cDNA FLJ11412 fs, clone HEMBA1000876. /FEA=mRNA /DB_XREF=gi:10432666 /UG=Hs.289094 Homo sapi 1.46
hypothetical protein NM_016449 1.46
recombination activating gene 1 NM_000448 1.46
regulated in glioma NM_006394 1.46
CD58 antigen, (lymphocyte function-associated antigen 3) BC005930 1.46
engulfment and cell motility 1 (ced-12 homolog, C. elegans) NM_014800 1.46
platelet activating receptor homolog NM_013308 1.46
hypothetical protein FLJ12650 NM_024522 1.46
POU domain, class 4, transcription factor 1 L20433 1.46
fracture callus 1 homolog (rat) NM_012192 1.46
cotector required for Sp1 transcriptional activation, subunit 3 (130kD) NM_004830 1.46
protein kinase, Y-linked NM_002760 1.46
aldehyde dehydrogenase 7 family, member A1 AU149534 1.46
hippocampal-like protein 4 BE550384 1.46
transient receptor potential channel 3 U47050 1.46
Cluster incl. AF070618:Homo sapiens clone 24627 mRNA sequence /cds=UNKNOWN /gb=AF070618 /gi=3283884 /ug=Hs.14074 /len=1179 1.46
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3 AJ251016 1.46
Consensus includes gb:AF203728.1 /DEF=Homo sapiens bone morphogenetic protein BMPY (BMPY) mRNA, partial cds. /FEA=mRNA /GEN=BMPY /PROD=bone morpho 1.46
cytochrome P450, subfamily IIIA (naphthalene oxidase), polypeptide 3 NM_000776 1.46
lipase protein BC001698 1.46
Consensus includes gb:AK023798.1 /DEF=Homo sapiens cDNA FLJ13736 fs, clone PLACE3000156. /FEA=mRNA /DB_XREF=gi:10435842 /UG=Hs.31532 Homo sapiens 1.46
Niemann-Pick disease, type C1 NM_000271 1.46
transcription factor 7-like 2 (T-cell specific, HMG-box) A1703074 1.46
Consensus includes gb:AL050388.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M2422 (from clone DKFZp564M2422); partial cds. /FEA=mRNA /GEN=DKFZp564M2422; 1.46
SRY (sex determining region Y-box 4) NM_003107 1.46
sodium channel, voltage-gated, type I, alpha polypeptide AF225985 1.46
retinoic acid- and interferon-inducible protein (58kD) NM_012420 1.46
LIM domain only 1 (hombolfin 1) NM_002315 1.46
folate hydrolase (prostate-specific membrane antigen) 1 NM_004478 1.46
cholinergic receptor, nicotinic, alpha polypeptide 3 K37981 1.46
ubiquitin 2 AK001029 1.46
hypothetical protein FLJ20813 NM_017961 1.46
MUF1 protein NM_006369 1.46
hypothetical protein MGC5384 NM_030972 1.46
lipase protein AA209239 1.46
ribose 5-phosphate isomerase A (ribose 5-phosphate epimerase) A1692341 1.46
KIAA0870 protein/actin NM_014977 1.46
HSKA-B protein AF070592 1.46
pregnancy-associated plasma protein-E AF311940 1.46

Consensus includes gb:Z84717.1 /DEF=Human DNA sequence from cDNA 16pHOG:19 from chromosome 16p13.3 /FEA=mRNA /DB_XREF=gi:1817571 /UG=Hs.305284 Z84717
 xylosyltransferase II NM_022167 1.46
 empty spiracles homolog 2 (Drosophila) AA78453 1.46
 PRO1693 protein AF118071 1.46
 potassium inwardly-rectifying channel, subfamily J, member 5 L47208 1.46
 hypothetical protein FLJ22684 NM_025048 1.46
 proprotein convertase subtilisin/kexin type 5 NM_008200 1.46
 hypothetical protein NM_018836 1.46
 B-cell CLL/lymphoma 2 M13594 1.46
 integral membrane protein 2A AL021786 1.46
 KIAA0664 protein AB014564 1.46
 BCL2-related ovarian killer NM_014204 1.46
 matrix metalloproteinase 16 (membrane-inserted) NM_022564 1.46
 crystallin, gamma D NM_008891 1.46
 ring finger protein 17 NM_031277 1.46
 Consensus includes gb:AU147851 /FEA=EST /DB_XREF=gi:1009372 /DB_XREF=est:AU147851 /CLONE=MAMMA1001878 /UG=Hs.183819 Homo sapiens cDNA FLJ12 AU147851 1.46
 integrin, beta 6 NM_000888 1.46
 hypothetical protein FLJ20241 AK023399 1.46
 spondin 1, (I-spondin) extracellular matrix protein A085290 1.46
 zinc finger protein 146 NM_007145 1.46
 hypothetical protein FLJ10783 NM_018217 1.46
 integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B) NM_000419 1.46
 Consensus includes gb:AV726900 /FEA=EST /DB_XREF=gi:1083632 /DB_XREF=est:AV726900 /CLONE=HTCABD05 /UG=Hs.7000 ESTs AV726900 1.46
 erythropoietin AF053356 1.46
 KIAA0644 gene product NM_014817 1.46
 protocadherin 1 (cadherin-like 1) AA461556 1.46
 BTG family, member 3 NM_008806 1.46
 v-rat simian leukemia viral oncogene homolog A (ras related) NM_005402 1.46
 hypothetical protein PRO2822 NM_018539 1.46
 inhibitor of growth family, member 1-like NM_001564 1.46
 nuclear antigen Sp100 NM_003113 1.46
 Consensus includes gb:AW968210 /FEA=EST /DB_XREF=gi:8158170 /DB_XREF=est:EST380405 /UG=Hs.293957 ESTs. Weakly similar to ALUC_HUMAN !!! ALU CLAS AW968210 1.46
 immunoglobulin lambda joining 3 AF234254 1.46
 T cell receptor delta locus X06557 1.46
 KIAA1356 protein AF352026 1.45
 immunoglobulin lambda-like polypeptide 2 L02326 1.45
 Consensus includes gb:AW970948 /FEA=EST /DB_XREF=gi:160793 /DB_XREF=est:EST383031 /UG=Hs.269403 ESTs AW970948 1.45
 solute carrier family 16 (monocarboxylic acid transporters), member 7 AF048608 1.45
 homolog of Xenopus Claspin NM_022111 1.45
 claudin 8 AL049977 1.45
 ecdonucleotide pyrophosphatase/phosphodiesterase 2 (autolysin) D49421 1.45
 protein with polyglutamine repeat: calcium (ca2+) homeostasis endoplasmic reticulum protein NM_004587 1.45
 Consensus includes gb:A088462 /FEA=EST /DB_XREF=gi:5591646 /DB_XREF=est:wn39e07.x1 /CLONE=IMAGE:2447844 /UG=Hs.238750 ESTs. Highly similar to serine A088462 1.45
 aldolase A, fructose-bisphosphate pseudogene 2 M21191 1.45
 carboxypeptidase B2 (plasma, carboxypeptidase U) NM_018413 1.45
 Consensus includes gb:U00956.1 /DEF=Human clone KDB5.1 (CAC)(n)(GTG)n repeat-containing mRNA. /FEA=mRNA /DB_XREF=gi:405058 /UG=Hs.104112 Human clone U00956 1.45
 Consensus includes gb:AL049328.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E026 (from clone DKFZp564E026). /FEA=mRNA /DB_XREF=gi:4500102 /UG=Hs.1356 AL049328 1.45
 plasmin 1 (I isoform) NM_002670 1.45
 potassium voltage-gated channel, Shal-related subfamily, member 3 AF120491 1.45
 hypothetical protein FLJ10815 AK022788 1.45
 putative peroxisome microbody protein 175.1 NM_015887 1.45
 KIAA1381 protein AK024376 1.45
 MAD, mothers against decapentaplegic homolog 8 (Drosophila) A028464 1.45
 PRO1398 protein NM_031289 1.45
 tight junction protein 1 (zona occludens 1) AA813018 1.45
 receptor-interacting serine-threonine kinase 2 AF027706 1.45
 cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 7 NM_000784 1.45
 T-box 6 AK022330 1.45
 ameloblastin, enamel matrix protein NM_016519 1.45
 AT2 receptor-interacting protein 1 AL098842 1.45
 CD3-epitope-associated protein; antisense to ERCC-1 NM_012099 1.45
 myristoylated alanine-rich protein kinase C substrate M68956 1.45

testicular hspidol expressed gene	NM_016585	1.45
phospholipase A2, group IID	AL158172	1.45
GATA binding protein 1 (globin transcription factor 1)	M30601	1.45
Ras homolog enriched in brain 2	D78132	1.45
hypothetical protein MGC13138	AA746280	1.45
ribosomal protein L18	AI660349	1.45
gb:NM_003723.1 /DEF=Homo sapiens caspase 13, apoptosis-related cysteine protease (CASP13), mRNA. /FEA=mRNA /GEN=CASP13 /PROD=caspase 13 /DB_XREF=	NM_003723	1.45
ribosomal protein S6 kinase, 90kD, polypeptide 6	NM_014496	1.45
mitogen-activated protein kinase 1 interacting protein 1	NM_021970	1.45
similar to slyltransferase 7 (alpha-N-acetylneuraminyl-2,3-betagalactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) E	NM_030965	1.45
pregnancy-induced growth inhibitor	NM_013370	1.45
hypothetical protein FLJ21901	NM_024622	1.45
tumor necrosis factor (ligand) superfamily, member 13	AF114011	1.45
hypothetical protein FLJ23263	NM_025115	1.45
Apobec-1 complementation factor; APOBEC-1 stimulating protein	NM_014576	1.45
CMRF35 leukocyte immunoglobulin-like receptor	NM_006678	1.45
serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antipapain, pigment epithelium derived factor), member 2	NM_000934	1.45
leukemia inhibitory factor receptor	NM_002310	1.45
Consensus includes gb:BF433429 /FEA=EST /DB_XREF=est:7q53a12.x1 /CLONE=IMAGE:3702046 /UG=Hs.289068 Homo sapiens cDNA FLJ11 BF433429		1.45
Consensus includes gb:U79265.1 /DEF=Human clone 23614 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710224 /UG=Hs.91375 Human clone 23614 mRNA sequence U79265		1.45
Consensus includes gb:U05004.1 /DEF=Homo sapiens mRNA; cDNA DKFZp568M0524 (from clone DKFZp568M0524). /FEA=mRNA /DB_XREF=gi:4884285 /UG=Hs.3C AL050043		1.45
Consensus includes gb:D29810.1 /DEF=Human mRNA for unknown product, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:704440 /UG=Hs.153445 Human mlf D29810		1.45
Rev/Rex activation domain binding protein-related	AF015042	1.45
Consensus includes gb:M16917 /DEF=Human glucocerebrosidase gene 10-2, 5 end /FEA=CDS_1 /DB_XREF=gi:183025 /UG=Hs.287396 Human glucocerebrosidase gene M16917		1.45
ZYG homolog	BE875849	1.45
lymphoprophthisis 1 (juvenile)	NM_000272	1.45
lymphoprophthisis alpha (TNF superfamily, member 1)	NM_000595	1.45
Consensus includes gb:A017119 /FEA=EST /DB_XREF=est:ov34a05.x1 /CLONE=IMAGE:1639184 /UG=Hs.144339 Human DNA sequence from c A017119		1.45
Consensus includes gb:AU143984 /FEA=EST /DB_XREF=est:AU143984 /CLONE=HEMBA1000568 /UG=Hs.301645 Homo sapiens cDNA FLJ100 AU143984		1.45
glutamate receptor, metabotropic 3	NM_000840	1.45
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson AB030075		1.45
Cluster incl. A937160/vp3f02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:2467443 /clone_end=3 /gb=A937160 /gb=5676030 /ug=Hs.7967 /len=552		1.45
senum constituent protein	NM_007061	1.45
nudix (nucleoside diphosphate linked moiety X)-type motif 7	AK028469	1.45
glycophorin A (includes MN blood group)	U00178	1.45
Consensus includes gb:AK024185.1 /DEF=Homo sapiens cDNA FLJ14123 fts, clone MAMMA1002155. /FEA=mRNA /DB_XREF=gi:10436502 /UG=Hs.269314 Homo sapiens cDNA AK024185		1.45
carbamoyl-phosphate synthetase 1, mitochondrial	AF154830	1.45
slatherin	NM_003154	1.45
gb:U05102.1 /DEF=Human IG processed VH-region mRNA, complete cds. /FEA=mRNA /GEN=IGH@ /PROD=immunoglobulin heavy chain V region /DB_XREF=gi:185528 /L05102		1.45
Consensus includes gb:AK025143.1 /DEF=Homo sapiens cDNA: FLJ21490 fts, clone COL05464. /FEA=mRNA /DB_XREF=gi:10437602 /UG=Hs.288700 Homo sapiens cDNA AK025143		1.44
postmeiotic segregation increased 2-like 6	D39503	1.44
steroidogenic acute regulatory protein	NM_000349	1.44
fer-1-like 3, myofibrin (C. elegans)	NM_013451	1.44
prostaglandin E synthase	NM_004878	1.44
Consensus includes gb:AL079289.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 35971. /FEA=mRNA /DB_XREF=gi:5102748 /UG=Hs.137154 /AL079289		1.44
hypothetical protein FLJ10675	NM_018176	1.44
Intersectin 1 (SH3 domain protein)	AF114488	1.44
serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1	NM_000602	1.44
lyyyl oxidase	AB028960	1.44
glutamate receptor, ionotropic, N-methyl D-aspartate 1	BE503425	1.44
lms-related tyrosine kinase 1 (vascular endothelial growth factor/vascular permeability factor receptor)	AF015731	1.44
killer cell lectin-like receptor subfamily D, member 1	AA058828	1.44
Consensus includes gb:AF052140.1 /DEF=Homo sapiens clone 23758 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360449 /UG=Hs.141055 Homo sapiens clone 23758 AF052140		1.44
Consensus includes gb:X63118 /DEF=H. sapiens DNA for oestrogen receptor gene promoter /FEA=CDS /DB_XREF=gi:35159 /UG=Hs.247938 H. sapiens DNA for oestrogen X63118		1.44
Mahlavu hepatocellular carcinoma	NM_006543	1.44
transcription factor AP-2 beta (activating enhancer binding protein 2 beta)	NM_003221	1.44
transmembrane 4 superfamily member (tetraspan NET-2)	NM_012338	1.44
hypothetical protein FLJ20950	NM_024952	1.44
Consensus includes gb:BF110551 /FEA=EST /DB_XREF=gi:10940241 /DB_XREF=est:7n38f08.x1 /CLONE=IMAGE:3567208 /UG=Hs.253600 ESTs, Highly similar to S50 BF110551		1.44
insulin-like growth factor binding protein 5	AW157548	1.44
lectin medin-2	NM_014921	1.44
solute carrier family 14 (urea transporter), member 1 (Kidd blood group)	NM_015865	1.44

KIAA1030 protein 1.44

EphA5 1.44

SA (rat hypertension-associated) homolog 1.44

ferrochelatase (protoporphyrin) 1.44

kinesin family member C3 1.44

JM1 protein 1.44

netrin 2-like (chicken) 1.44

hepatocellular carcinoma novel gene-3 protein 1.44

hypothetical protein FLJ23554 1.44

solute carrier family 4, sodium bicarbonate cotransporter, member 4 1.44

ets variant gene 6 (TEL oncogene) 1.44

hypothetical protein PRO1895 1.44

hypothetical protein FLJ20337 1.44

splicing factor, arginine/serine-rich 11 1.44

phosphatidylinositol glycan, class K 1.44

hypothetical protein FLJ10786 1.44

B-cell CLL/lymphoma 11A (zinc finger protein) 1.44

empty spiracles homolog 1 (Drosophila) 1.44

mucin 5, subtype B, tracheobronchial 1.44

holocytochrome c synthase (cytochrome c heme-lyase) 1.44

Consensus includes gb:AW974995 /FEA=EST /DB_XREF=estEST387100 /UG=Hs.292787 ESTs

KIAA0699 protein 1.44

KIAA1922 protein 1.44

killer cell lectin-like receptor subfamily D, member 1 1.44

forhead box E1 (thyroid transcription factor 2) 1.44

hypothetical protein FLJ22031 1.44

Consensus includes gb:AF090910.1 /DEF=Homo sapiens clone HQ0262 /FEA=mrna /DB_XREF=gi:5690195 /UG=Hs.236760 Homo sapiens clone HQ0262

Consensus includes gb:AK026439.1 /DEF=Homo sapiens cDNA: FLJ22786 fls, clone KIAA2150, /FEA=mrna /DB_XREF=gi:10439304 /UG=Hs.306848 Homo sapiens cDNA: AK026439

kalikrein 14 1.44

ankyrin 1, erythrocytic 1.44

ATPase, Ca++ transporting, plasma membrane 3 1.44

NCF 2 cell line derived transforming sequence 1.44

synapsin I 1.44

hypothetical protein FLJ20712 1.44

DiGeorge syndrome critical region gene 2 1.44

RAD54-like (S. cerevisiae) 1.44

ret finger protein-like 3 1.44

phosphodiesterase 1A, calmodulin-dependent 1.44

STAT induced STAT inhibitor-2 1.44

glutamate receptor, ionotropic, N-methyl D-aspartate 1 1.44

hypothetical protein FLJ12331 1.44

cytochrome P450, subfamily IVF, polypeptide 8 1.44

adaptor-related protein complex 3, sigma 1 subunit 1.44

proprotein convertase subtilisin/kexin type 2 1.44

nuclear receptor subfamily 5, group A, member 1 1.44

neuronal thread protein 1.44

solute carrier family 21 (organic anion transporter), member 3 1.44

disabled homolog 1 (Drosophila) 1.44

microtubule-associated protein 1A 1.44

matrix metalloproteinase 17 (membrane-inserted) 1.44

homeo box A3 1.44

heterogeneous nuclear ribonucleoprotein H1 (H) 1.44

glycogen synthase 2 (liver) 1.44

NGF-A binding protein 1 (EGR1 binding protein 1) 1.44

Fc fragment of IgG, high affinity Ia, receptor for (CD64) 1.44

UDP-Gal4betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 2 1.44

phosphodiesterase 3A, cGMP-inhibited 1.44

hypothetical protein MGC4643 1.44

anaplastic lymphoma kinase (Kt-1) 1.44

adenylyate cyclase 9 1.44

aggrekan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122) 1.44

Zc family member 1 (odd-paired homolog, Drosophila) 1.44

fms-related tyrosine kinase 1 (vascular endothelial growth factor/vascular permeability factor receptor)
 MCH3a minichromosome maintenance deficient 3 (S. cerevisiae) associated protein
 homolog of yeast ATP12
 Consensus includes gb:AK022038.1 /DEF=Homo sapiens cDNA FLJ11976 fls, done HEMBB1001253. /FEA=mRNA /DB_XREF=gi:10433356 /UG=Hs.296679 Homo sapie
 KIAA0063 gene product
 hypothetical protein R32184_1
 solute carrier family 5 (neutral amino acid transporters, system A), member 4
 phosphoinositide-3-kinase, catalytic, beta polypeptide
 eve, even-skipped homeo box homolog 1 (Drosophila)
 RAD51-like 1 (S. cerevisiae)
 KIAA0318 protein
 Consensus includes gb:S80491.1 /DEF=stem cell factor (alternatively spliced) human, preimplantation embryos, blastocysts, mRNA Partial, 180 nt. /FEA=mRNA /GEN=ste
 cadherin 15, M-cadherin (myotubule)
 T-box 1
 Consensus includes gb:AK024393.1 /DEF=Homo sapiens cDNA FLJ14331 fls, done PLACE4000320. /FEA=mRNA /DB_XREF=gi:10436771 /UG=Hs.288569 Homo sapie
 KIAA0977 protein
 tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain
 solute carrier family 18 (folate transporter), member 1
 mitochondrial ribosomal protein L33
 Consensus includes gb:AL035295.1 /DEF=H. sapiens gene from PAC 106H8, similar to Dynamin. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:4200217 /UG=Hs
 hypothetical protein DKFZ0761K1824
 likely homolog of mouse heparin/heparin sulfate-glucuronic acid C5-epimerase
 hypothetical protein DKFZp761K1423
 adenylate cyclase 2 (brain)
 Consensus includes gb:AK024355.1 /DEF=Homo sapiens cDNA FLJ14293 fls, done PLACE1007868. /FEA=mRNA /DB_XREF=gi:10436719 /UG=Hs.141009 Homo sapie
 hypothetical protein OVC10-2
 hypothetical protein FLJ13941
 KIAA0672 gene product
 myeloid cell nuclear differentiation antigen
 melanoma cell adhesion molecule
 Lin-7b protein; likely ortholog of mouse LIN-7B; mammalian LIN-7 protein 2
 nuclear fragile X mental retardation protein interacting protein 1
 HSPC048 protein
 Consensus includes gb:AK023187.1 /DEF=Homo sapiens cDNA FLJ13125 fls, done NT2RP3002877. /FEA=mRNA /DB_XREF=gi:10435001 /UG=Hs.287551 Homo sapie
 chloride channel 1, skeletal muscle (Thomsen disease, autosomal dominant)
 TIA1 cytotoxic granule-associated RNA binding protein
 zinc finger protein 108
 calmodulin 2 (phosphorylase kinase, delta)
 mannosyl (alpha-1,3)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme B
 KIAA0485 protein
 Consensus includes gb:AF052173.1 /DEF=Homo sapiens clone 24417 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360484 /UG=Hs.306253 Homo sapiens clone 24417 AF052173
 Consensus includes gb:BF002254. /FEA=EST /DB_XREF=gi:10702529 /DB_XREF=est:7h02a03.x1 /CLONE=IMAGE.3314764 /UG=Hs.143600 type II Golgi membrane pro
 KIAA0680 gene product
 midline 1 (Optiz/BBB syndrome)
 B-cell CLL/lymphoma 10
 integrin, beta 3 (platelet glycoprotein IIb, antigen CD61)
 discs, large (Drosophila) homolog 2 (chapsyn-110)
 hypothetical protein FLJ11043
 Consensus includes gb:N64802 /FEA=EST /DB_XREF=gi:1212631 /DB_XREF=est:y231b05.s1 /CLONE=IMAGE.284625 /UG=Hs.173684 Homo sapiens mRNA; cDNA DK1 N54802
 Consensus includes gb:AI206888 /FEA=EST /DB_XREF=gi:3765560 /DB_XREF=est:q35a04.x1 /CLONE=IMAGE.1942830 /UG=Hs.154131 ESTs
 tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)
 hypothetical protein FLJ22477
 D15F37 (pseudogene)
 Consensus includes gb:AL049442.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N1720 (from done DKFZp586N1720). /FEA=mRNA /DB_XREF=gi:4500222 /UG=Hs.71
 phosphatidylinositol-4-phosphate 5-kinase, type I, beta
 myeloid/lymphoid or mixed-lineage leukemia (rithorax homolog, Drosophila); translocated to, 2
 hypothetical protein FLJ20105
 four and a half LIM domains 1
 hypothetical protein FLJ11113
 disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)
 host cell factor C1 (VP16-accessory protein)
 MD-2 protein
 lethal giant larvae homolog 1 (Drosophila)

allograft inflammatory factor 1
 olfactory receptor, family 1, subfamily K, member 1
 arylsulfatase A
 hematopoietic PBX-interacting protein
 glycoprotein M6B
 hypothetical protein PRO1728
 Consensus includes gb:L37198.1 /DEF=Homo sapiens (clone B3B3E13) Huntingtons disease candidate region mRNA fragment. /FEA=mRNA /DB_XREF=gi:600519 /UG=L37198
 dickkopf homolog 3 (Xenopus laevis)
 Consensus includes gb:AK026667.1 /DEF=Homo sapiens cDNA: FLJ23014 fis, clone LNG00806. /FEA=mRNA /DB_XREF=gi:10439568 /UG=Hs.306862 Homo sapiens cDNA: AK026667
 hypothetical protein FLJ10761
 Consensus includes gb:AK022442.1 /DEF=Homo sapiens cDNA FLJ12380 fis, clone MAMMA1002556. /FEA=mRNA /DB_XREF=gi:10433838 /UG=Hs.300934 Homo sapiens cDNA: AK022442
 v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)
 forhead box L1
 synaptophysin, gamma 2
 DLEU2L
 chromosome 4 open reading frame 5
 Consensus includes gb:AL110227.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434J194 (from clone DKFZp434J194). /FEA=mRNA /DB_XREF=gi:10433838 /UG=Hs.300934 Homo sapiens cDNA: AK022442
 KIAA1305 protein
 Consensus includes gb:AL030995 /DEF=Human DNA sequence from clone 1170D6 on chromosome Xq22.3-23. Contains a pseudogene similar to U-SNRNP. associated (AL030995)
 Notch domain and RLD 3
 KIAA0257 protein
 Consensus includes gb:AL157902 /DEF=Human DNA sequence from clone RPA4-675C20 on chromosome 1p13.2. Contains the 3' end of the MAN1A2 gene for mannosidase AL157902
 major histocompatibility complex, class II, DP alpha 1
 X17013 B subtilis lys gene for dimethylamine decarboxylase corresponding to nucleotides 350-1345 of X17013 (-5, -M, -3 represent transcript regions 5 prime, Middle, ar X17013
 betacellulin
 Consensus includes gb:AL133396 /DEF=Human DNA sequence from clone RPS-1068H6 on chromosome 20q11.1-11.23. Contains a pseudogene similar to ID11 (isopentyl) AW675655
 Consensus includes gb:AW675655 /FEA=EST /DB_XREF=gi:7540890 /DB_XREF=est:ba52e01.x1 /CLONE=IMAGE:2900184 /UG=Hs.314158 ESTs
 nucleoporin 62KD
 TAF₉-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD
 undifferentiated embryonic cell transcription factor 1
 Consensus includes gb:AL035535 /DEF=Human DNA sequence from clone 217P22 on chromosome 6p21.1-21.31. Contains part of a novel Dynein heavy chain family prot AL035535
 retinoblastoma binding protein 2
 ATP-binding cassette, sub-family A (ABC1), member 3
 Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)
 calcium channel, voltage-dependent, alpha 2/delta subunit 2
 radixin
 calcium/calmodulin-dependent protein kinase (CaM kinase) II alpha
 hypothetical protein, estradiol-induced
 galactosidase, alpha
 peroxisomal biogenesis factor 3
 serine racemase
 lipin 1
 hypothetical protein FLJ20069
 Consensus includes gb:AL133018.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434F0327 (from clone DKFZp434F0327). /FEA=mRNA /DB_XREF=gi:6453498 /UG=Hs.271 /AL133018
 small optic tectal homolog (Drosophila)
 Consensus includes gb:BE658194 /FEA=EST /DB_XREF=gi:10372818 /DB_XREF=est:7g19c05.x1 /CLONE=IMAGE:3306920 /UG=Hs.323053 Homo sapiens mRNA full length BE658194
 potassium inwardly-rectifying channel, subfamily J, member 14
 phosphodiesterase 7B
 hyaluronoglucosaminidase 1
 brain-specific angiogenesis inhibitor 3
 HT001 protein
 ankyrin repeat domain 2 (stretch responsive muscle)
 calcium binding protein 1 (calbrain)
 forhead box C2 (MFX-1, mesenchyme forhead 1)
 hypothetical protein DKFZp781B2423
 heat shock transcription factor 2 binding protein
 Consensus includes gb:AA78300 /FEA=EST /DB_XREF=gi:4371526 /DB_XREF=est:tm39e01.x1 /CLONE=IMAGE:2160504 /UG=Hs.192789 ESTs, Weakly similar to ALUK AA78300
 Dlx-binding protein IDAX (inhibition of the Dlx and Axin complex)
 ATPase, Ca++ transporting, type 2C, member 1
 deleted in lymphocytic leukemia, 2
 solute carrier family 6 (neurotransmitter transporter, glycine), member 5
 phosphodiesterase 4D Interacting protein (myomegalin)

wingless-type MMTV integration site family, member 108
hypothetical protein FLJ22595
Consensus includes gb:AL110248.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N021 (from clone DKFZp434N021) /FEA=mRNA /DB_XREF=gi:5817198 /UG=Hs.2087 AL110248 1.42
protocadherin beta 11
epsin 3
haptoglobin-related protein
alpha-catenin-like protein
cell cycle progression 8 protein
tachykinin, precursor 1 (substance K, neurokinin 1, neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide gamma)
paired box gene 3 (Wardenburg syndrome 1)
hypothetical protein FLJ14639
arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)
fetuin B
ras-like protein
hypothetical protein
hypothetical protein FLJ13576
tumor necrosis factor (ligand) superfamily, member 12
Consensus includes gb:AL049430.1 /DEF=Homo sapiens mRNA; cDNA DKFZp568H201 (from clone DKFZp568H201) /FEA=mRNA /DB_XREF=gi:4500210 /UG=Hs.3020 AL049430 1.42
ectodermal dysplasia 1, anhidrotic
ring finger protein 4
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calciuretin A gamma)
serine protease inhibitor, Kunitz type, 3
hypothetical protein FLJ11100
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4
S164 protein
Consensus includes gb:AK024445.1 /DEF=Homo sapiens mRNA for FLJ00035 protein, partial cds. /FEA=mRNA /GEN=FLJ00035 /PROD=FLJ00035 protein /DB_XREF=gi:AK024445 1.42
stailidase 3 (membrane stailidase)
protein kinase, cGMP-dependent, type II
opioid receptor, mu 1
E2F transcription factor 5, p130-binding
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2
Consensus includes gb:AW450360 /FEA=EST /DB_XREF=gi:6991136 /DB_XREF=est:U1-H-B13-akn-c-11-U1.s1 /CLONE=IMAGE:2734845 /UG=Hs.98834 ESTs 1.42
KIAA0161 gene product
translocase of inner mitochondrial membrane 17 homolog A (yeast)
t-complex 1
Consensus includes gb:BE880245 /FEA=EST /DB_XREF=gi:10329021 /DB_XREF=est:601490812F1 /CLONE=IMAGE:3893018 /UG=Hs.164038 Homo sapiens AKAP3501 BE880245 1.42
transcription elongation factor B (SIII), polypeptide 2 (18kD, elongin B)
dual specificity phosphatase 4
GTP-binding protein Rho7
kinesin family member 13A
lengsin
pregnancy specific beta-1-glycoprotein 3
mitochondrial ribosomal protein 63
Consensus includes gb:AK024128.1 /DEF=Homo sapiens cDNA FLJ14086 fs, clone HEMBB1001197 /FEA=mRNA /DB_XREF=gi:10436433 /UG=Hs.306865 Homo sapie AK024128 1.42
neuregulin 1
LIV-1 protein, estrogen regulated
cadherin-like protein VR20
secretogranin III
neurogenin 1
nucleobindin 1
ets variant gene 1
hypothetical protein FLJ12270
glutamate receptor, ionotropic, kainate 5
hypothetical protein
GTP binding protein 1
Usher syndrome 2A (autosomal recessive, mild)
gap junction protein, alpha 8, 50kD (connexin 50)
gq9qj-67
Consensus includes gb:AL049331.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564I066 (from clone DKFZp564I066) /FEA=mRNA /DB_XREF=gi:4500106 /UG=Hs.207801 AL049331 1.42
hypothetical protein FLJ20069
S100 calcium binding protein A6 (calycidin)
kinesin-like 3

acyltransferase (neutrophil)
 Consensus includes gb:BE742268 /FEA=EST /DB_XREF=gi:10156260 /DB_XREF=est:601575686F.1 /CLONE=IMAGE:3836709 /UG=Hs.281706 sortilin 1
 Consensus includes gb:R43279 /FEA=EST /DB_XREF=gi:821386 /DB_XREF=est:yg17c12.s1 /CLONE=IMAGE:32731 /UG=Hs.22574 ESTs
 hypothetical protein FLJ11800
 Consensus includes gb:AA156865 /FEA=EST /DB_XREF=gi:1728460 /DB_XREF=est:z20008.s1 /CLONE=IMAGE:502503 /UG=Hs.22370 Homo sapiens mRNA; cDNA DK
 DKFZP434N043 protein
 cyclin T2
 p8 protein (candidate of metastasis 1)
 bone morphogenetic protein 2
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
 cyclochrome b-245, beta polypeptide (chronic granulomatous disease)
 testis intracellular mediator protein
 toll-like 2
 Consensus includes gb:AL139318 /DEF=Human DNA sequence from clone RP11-124B17 on chromosome 13. Contains the DCT gene for dopachrome tautomerase (dopa
 Consensus includes gb:AK024945.1 /DEF=Homo sapiens cDNA; FLJ21292 fls, clone COL01869 /FEA=mRNA /DB_XREF=gi:10437367 /UG=Hs.306732 Homo sapiens cDNA
 adenylylase 3
 UDP-N-acetyl-alpha-D-galactosamine-4-epimerase (N-acetylneuraminyl)-galactosylglucosyltransferase (GALNAc-T)
 transmembrane 7 superfamily member 1 (upregulated in kidney)
 guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF
 PRO0038 protein
 guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF
 protein tyrosine phosphatase, receptor type, M
 cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)
 phospholipase A2, group V
 nucleoporin 153KO
 parathyroid hormone receptor 1
 chitinase, di-N-acetyl-
 glial cells missing homolog b (Drosophila)
 RAP1, GTPase activating protein 1
 PRO0457 protein
 malic enzyme 1, NAD(P)+-dependent, cytosolic
 Consensus includes gb:U06813.1 /DEF=Homo sapiens serotonin-7 receptor pseudogene, complete sequence. /FEA=mRNA /PROD=serotonin-7 receptor /DB_XREF=gi:31 U68813
 Consensus includes gb:AL571723 /FEA=EST /DB_XREF=gi:12929303 /DB_XREF=est:AL571723 /CLONE=CS01031YD10 (3 prime) /UG=Hs.7247 ESTs, Weakly similar 1
 vesicle-associated membrane protein 3 (cellubrevin)
 tropomyosin 2 (beta)
 G protein-coupled receptor kinase 2-like (Drosophila)
 chemokine (C-C motif) receptor 3
 hypothetical protein MGC2718
 hypothetical protein FLJ20071
 1,2-alpha-mannosidase IC
 homolog of mouse quaking QKI (KH domain RNA binding protein)
 chromosome 1 open reading frame 24
 ichthy homolog E3 ubiquitin protein ligase (mouse)
 Consensus includes gb:AK024860.1 /DEF=Homo sapiens cDNA; FLJ21207 fls, clone COL00362. /FEA=mRNA /DB_XREF=gi:10437268 /UG=Hs.306708 Homo sapiens cDNA
 KIAA0443 gene product
 KIAA0831 protein
 hypothetical protein FLJ23510
 heat shock 70kD protein 1A
 interleukin 18 (lymphocyte chemoattractant) factor
 Consensus includes gb:AF05011.1 /DEF=Homo sapiens clone 24587 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005734 /UG=Hs.137173 Homo sapiens clone 24587
 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 1
 FOS-like antigen 2
 collagen, type XI, alpha 1
 transcription elongation factor A (SII)-like 1
 Consensus includes gb:AL049437.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566E1120 (from clone DKFZp566E1120) /FEA=mRNA /DB_XREF=gi:4500220 /UG=Hs.10
 testis expressed sequence 11
 adaptor-related complex 1, mu 2 subunit
 phospholipase A2, group VI (cytosolic, calcium-independent)
 lymphocyte antigen 84 homolog, activating NK-receptor; NK-p46, (mouse)
 Consensus includes gb:AL110180.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J2118 (from clone DKFZp564J2118) /FEA=mRNA /DB_XREF=gi:5817104 /UG=Hs.306
 ras homolog gene family, member 1
 hypothetical protein FLJ10803
 WD repeat domain 9

laminin, alpha 3 (nicotin (150KD), kalinin (165KD), BM600 (150KD), epiegtrin) NM_000227 1.42

BTG family, member 3 A1765445 1.42

fatty acid binding protein 7, brain NM_001446 1.42

hypothetical protein FLJ20401 NM_017805 1.41

decidual protein induced by progesterone AL136653 1.41

myristoylated alanine-rich protein kinase C substrate NM_002356 1.41

Consensus includes gb:AL133075.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M0317.1 (from done DKFZp434M0317). /FEA=mRNA /DB_XREF=gi:6453518 /UG=Hs.27 AF133075 1.41

gb:AF130051.1 /DEF=Homo sapiens clone FLB3535 PRO0898 mRNA, complete cds. /FEA=mRNA /PROD=PRO0898 /DB_XREF=gi:11493408 /UG=Hs.306860 Homo sap AF130051 1.41

Cytokeratin 20 W58342 1.41

hypothetical protein CG003 NM_007354 1.41

pustative GR6 protein BE565369 1.41

coagulation factor II (thrombin) receptor-like 1 NM_001251 1.41

CD68 antigen NM_007374 1.41

sine oculis homeobox homolog 6 (Drosophila U19713 1.41

allograft inflammatory factor 1 U53823 1.41

occludin BC000039 1.41

hypothetical protein NM_018162 1.41

hypothetical protein FLJ10833 NM_017416 1.41

interleukin 1 receptor accessory protein-like 2 AF028219 1.41

deleted in liver cancer 1 NM_000231 1.41

sarcoglycan, gamma (35KD dystrophin-associated glycoprotein) NM_013938 1.41

olfactory receptor, family 10, subfamily H, member 3 NM_002002 1.41

Fc fragment of IgC, low affinity II, receptor for (CD23A) NM_004183 1.41

Consensus includes gb:AF009287.1 /DEF=Homo sapiens clone FBA1 Cn-du-chat region mRNA. /FEA=mRNA /DB_XREF=gi:2331089 /UG=Hs.102238 Homo sapiens dbn AF009287 1.41

vitelliform macular dystrophy (Best disease, bestrophin) NM_005213 1.41

cyretatin A (steflin A) NM_004204 1.41

phosphatidylinositol glycan, class O NM_005839 1.41

Consensus includes gb:AW085172 /FEA=EST /DB_XREF=gi:6040324 /DB_XREF=est:xe06g09.x1 /CLONE=IMAGE:2606368 /UG=Hs.243366 ESTs, Highly similar to KPCI AW085172 1.41

serine/arginine repetitive matrix 1 NM_004204 1.41

Consensus includes gb:AU148024 /FEA=EST /DB_XREF=gi:11009545 /DB_XREF=est:AU148024 /CLONE=IMAGE:1002392 /UG=Hs.302075 Homo sapiens cDNA FLJ12 AU148024 1.41

forthead box O1A (maodomyosarcoma) NM_002280 1.41

Consensus includes gb:AU148024 /FEA=EST /DB_XREF=gi:11009545 /DB_XREF=est:AU148024 /CLONE=IMAGE:1002392 /UG=Hs.302075 Homo sapiens cDNA FLJ12 AU148024 1.41

Consensus includes gb:AK021930.1 /DEF=Homo sapiens cDNA FLJ11868 fls, clone HEMBA1008993. /FEA=mRNA /DB_XREF=gi:10433231 /UG=Hs.302280 Homo saple AK021930 1.41

KIAA1032 protein BE783668 1.41

Consensus includes gb:L13283.1 /DEF=Homo sapiens (clone MG2-5-12) mucin (MG2) mRNA, complete polyA site. /FEA=mRNA /DB_XREF=gi:292518 /UG=Hs.103944 H.L13283 1.41

FAT tumor suppressor homolog 1 (Drosophila) NM_005245 1.41

Consensus includes gb:AK46341 /FEA=EST /DB_XREF=gi:4083347 /DB_XREF=est:qp50b09.x1 /CLONE=IMAGE:1926425 /UG=Hs.298998 Homo sapiens, Similar to hyp AK46341 1.41

CD36 antigen (collagen type I receptor, thrombospondin receptor) M88399 1.41

RCE1 homolog, prenyl protein protease (S. cerevisiae) AF121951 1.41

retinoid acid receptor-beta associated open reading frame NM_015854 1.41

Consensus includes gb:AK024568.1 /DEF=Homo sapiens cDNA: FLJ20915 fls, clone ADSE00692. /FEA=mRNA /DB_XREF=gi:10436878 /UG=Hs.306691 Homo sapiens c AK024568 1.41

hypothetical protein FLJ11282 AK023539 1.41

neurtin NM_016588 1.41

ectonucleic translation initiation factor 3, subunit 5 (epsilon, 47KD) AL001896 1.41

Consensus includes gb:AU148187 /FEA=EST /DB_XREF=gi:5110251 /DB_XREF=est:wp48b05.x1 /CLONE=IMAGE:2368305 /UG=Hs.194051 Homo sapiens mRNA; cDNA AU148187 1.41

gb:AF118715.1 /DEF=Homo sapiens PRO2829 mRNA, complete cds. /FEA=mRNA /PROD=PRO2829 /DB_XREF=gi:7899928 /UG=Hs.236256 Homo sapiens PRO2829 m AF118715 1.41

Consensus includes gb:AK026847.1 /DEF=Homo sapiens cDNA: FLJ23194 fls, clone REC00490. /FEA=mRNA /DB_XREF=gi:10439802 /UG=Hs.306887 Homo sapiens c AK026847 1.41

KIAA0830 protein AL573201 1.41

guanine nucleotide binding protein (G protein), alpha (inhibiting activity polypeptide 1) AL049933 1.41

Consensus includes gb:AW771015 /FEA=EST /DB_XREF=gi:7703066 /DB_XREF=est:hms5q005.x1 /CLONE=IMAGE:3027081 /UG=Hs.279020 Homo sapiens mRNA; cDN AW771015 1.41

gb:NM_022125.1 /DEF=Homo sapiens immortalization-upregulated protein (IMUP), mRNA. /FEA=mRNA /GEN=IMUP /PROD=immortalization-upregulated protein /DB_XRI NM_022125 1.41

thrombospondin 1 BE589867 1.41

hypothetical protein FLJ13110 BE535746 1.41

Consensus includes gb:X69383 /DEF=H sapiens gene for T cell receptor gamma V region 5 /FEA=CDS_2 /DB_XREF=gi:37009 /UG=Hs.247801 H.sapiens gene for T cell r X69383 1.41

sushi-repeat-containing protein, X chromosome NM_005307 1.41

gb:NM_005179.1 /DEF=Homo sapiens avian lymphoma virus-derived transforming sequence (BLYM), mRNA. /FEA=CDS /GEN=BLYM /PROD=avian lymphoma virus-deri NM_005179 1.41

ADP-ribosylation factor 4-like NM_001661 1.41

Consensus includes gb:BF109941 /FEA=EST /DB_XREF=gi:10939631 /DB_XREF=est:772005.x1 /CLONE=IMAGE:3527024 /UG=Hs.27183 Homo sapiens clone 23940 m BF109941 1.41

Rho guanine exchange factor (GEF) 15 AK023853 1.41

cadherin 5, type 2, VE-cadherin (vascular epithelium) NM_001795 1.41

CD33 antigen (gp87) NM_001772 1.41

hypothetical protein FLJ13385 NM_024853 1.41

cholinergic receptor, nicotinic, alpha polypeptide 3 BC000513 1.41

KIAA0665 gene product AW243089 1.41

hypothetical protein
 peroxisomal biogenesis factor 3
 contactin associated protein-like 2
 dual oxidase 2
 transcriptional adaptor 3 (ADA3, yeast homolog-like (PCAF histone acetylase complex)
 Consensus includes gb:AL117652.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1324 (from done DKFZp586B1324). /FEA=mRNA /DB_XREF=gi:5912244 /UG=Hs.271 AL117652
 tropotin T1, skeletal, slow
 immunoglobulin heavy constant mu
 Consensus includes gb:AV1718487 /FEA=EST /DB_XREF=gi:10815639 /DB_XREF=est:AV1718487 /CLONE=GLCFVH09 /UG=Hs.282656 ESTs
 Consensus includes gb:AU679024 /FEA=EST /DB_XREF=gi:4893206 /DB_XREF=est:u71a10.x1 /CLONE=IMAGE:2256474 /UG=Hs.201420 ESTs, Moderately similar to AL1679024
 caudal type homeo box transcription factor 2
 KIAA1000 protein
 zinc finger protein 174
 putative zinc finger protein NY-REN-34 antigen
 chromosome 8 open reading frame 2
 protocadherin gamma subfamily A, 8
 anysaufatase F
 zinc finger protein 287
 LIM domain binding 2
 similar to MYOSIN HEAVY CHAIN, CARDIAC MUSCLE ALPHA ISOFORM (MYHC-ALPHA) (M. musculus)
 pyruvate dehydrogenase (liposimile) alpha 2
 Consensus includes gb:R37655 /FEA=EST /DB_XREF=gi:795111 /DB_XREF=est:y61403.s1 /CLONE=IMAGE:26881 /UG=Hs.194761 H. sapiens IMAGE cDNA clone 2688 R37655
 RNA binding motif protein 9
 piccolo (presynaptic cytomatrix protein)
 KIAA0485 protein
 collagen, type I, alpha 1
 golgi phosphoprotein 1
 KIAA0030 protein
 Consensus includes gb:AU742810 /FEA=EST /DB_XREF=gi:5111098 /DB_XREF=est:wq46a05.x1 /CLONE=IMAGE:2368160 /UG=Hs.131057 ESTs, Moderately similar to C AU742810
 gb:AF116680.1 /DEF=Homo sapiens PRO2168 mRNA, complete cds. /FEA=mRNA /PROD=PRO2168 /DB_XREF=gi:7959876 /UG=Hs.249715 Homo sapiens PRO2168 m AF116689
 POM (POM121 rat homolog) and ZP3 fusion
 hypothetical protein MGC5540
 protein kinase, AMP-activated, gamma 2 non-catalytic subunit
 steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1)
 Kruppel-type zinc finger (C2H2)
 Consensus includes gb:AF070571.1 /DEF=Homo sapiens clone 24739 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387940 /UG=Hs.44690 Homo sapiens clone 24739 AF070571
 potassium large conductance calcium-activated channel, subfamily M, beta member 4
 Consensus includes gb:AW301775 /FEA=EST /DB_XREF=gi:8711452 /DB_XREF=est:w63a06.x1 /CLONE=IMAGE:2766754 /UG=Hs.254727 ESTs
 Consensus includes gb:AW301775 /FEA=EST /DB_XREF=gi:10739541 /DB_XREF=gi:10739541 /DB_XREF=est:601557983F1 /CLONE=IMAGE:3827709 /UG=Hs.2631 desmoglein 2 /FL=gb-NM_020427
 ARS component B
 heterogenous nuclear ribonucleoprotein U (scaffold attachment factor A)
 sialyltransferase 4C (beta-galactosidase alpha-2,3-sialyltransferase)
 sine oculis homeobox homolog 1 (Drosophila)
 basic transcription element binding protein 1
 Consensus includes gb:AF088790.1 /DEF=Homo sapiens aconitase precursor (ACON) mRNA, nuclear gene encoding mitochondrial protein, partial cds. /FEA=mRNA /GEI AF088790
 Consensus includes gb:S83374.1 /DEF=glutamate transporter II variant BHGT II (5 region) human, brain and spinal cord, mRNA Partial Mutant, 129 nt. /FEA=mrna /DI S83374
 Consensus includes gb:T87235 /FEA=EST /DB_XREF=gi:715577 /DB_XREF=est:ydi1106.s1 /CLONE=IMAGE:22392 /UG=Hs.13223 Homo sapiens mRNA full length inser T87235
 DNA segment on chromosome X (unique) 9878 expressed sequence
 hypothetical protein FLJ13842
 G protein-coupled receptor 37 (endothelin receptor type B-like)
 membrane protein, palmitoylated 1 (55kd)
 Consensus includes gb:AU146952 /FEA=EST /DB_XREF=gi:11008473 /DB_XREF=est:AU146952 /CLONE=HEMBA1001962 /UG=Hs.306624 Homo sapiens cDNA FLJ121 AU146952
 nuclear receptor subfamily 4, group A, member 3
 Consensus includes gb:AU146952 /FEA=EST /DB_XREF=gi:11008473 /DB_XREF=est:AU146952 /CLONE=HEMBA1001962 /UG=Hs.306624 Homo sapiens cDNA FLJ121 AU146952
 KIAA0205 gene product
 fibroblast growth factor receptor 3 (MAGUK p55 subfamily member 3)
 leptin receptor
 hypothetical protein FLJ10547

RAS protein activator like 2
Consensus includes gb:AK024081.1 /DEF=Homo sapiens cDNA FLJ14019 fls. clone HEMBA1002503. /FEA=mRNA /DB_XREF=gi:10436370 /UG=Hs.287613 Homo sapiens KIAA0768 protein
Consensus includes gb:AL137312.1 /DEF=Homo sapiens mRNA; cDNA DKFZp781K23121 (from clone DKFZp781K23121). /FEA=mRNA /DB_XREF=gi:6807797 /UG=Hs. AL137312
serine/threonine kinase 3 (STE20 homolog, yeast)
gb:NM_022155.1 /DEF=Homo sapiens PP3227 protein (PP3227), mRNA. /FEA=mRNA /GEN=PP3227 /PROD=PP3227 protein /DB_XREF=gi:11545901 /UG=Hs.302027 F NM_022155
ventral anterior homeobox 2
Integrin, beta 8
gallin
CASP8 and FADD-like apoptosis regulator
hypothetical protein FLJ11168
carcinoembryonic antigen-related cell adhesion molecule 5
DKFZP584C103 protein
Consensus includes gb:AV704232 /FEA=EST /DB_XREF=gi:10721550 /DB_XREF=est:AV704232 /CLONE=ADBBLB11 /UG=Hs.228096 Homo sapiens cDNA FLJ11682 fls. AV704232
hypothetical protein
phosphoserine phosphatase
hepatocyte nuclear factor 4, gamma
collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital)
matrix metalloproteinase 28
tripartite motif-containing 14
hypothetical protein MGC2652
fasciculation and elongation protein zeta 1 (zyg11)
hypothetical protein FKSG25
low density lipoprotein receptor-related protein 4
B-cell CLL/lymphoma 2
CGI-109 protein
retinoid X receptor, beta
KIAA0923 protein
hypothetical protein FLJ10986
B9 protein
hypothetical protein FLJ14146
GDP-ucose transporter 1
Consensus includes gb:AK024129.1 /DEF=Homo sapiens cDNA FLJ14087 fls. clone HEMBB1001315. /FEA=mRNA /DB_XREF=gi:10436434 /UG=Hs.287619 Homo sapiens AK024129
hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1
Consensus includes gb:BE378994 /FEA=EST /DB_XREF=gi:9324359 /DB_XREF=est:6012362471 /CLONE=IMAGE:3608626 /UG=Hs.271717 ESTs. Weakly similar to AL BE378994
arginine vasopressin (neurophysin II, antidiuretic hormone, diabetes insipidus, neurohypophyseal)
Consensus includes gb:X9637.1 /DEF=H.sapiens mRNA sequence (16p11.2). /FEA=mRNA /DB_XREF=gi:29573 /UG=Hs.307208 H.sapiens mRNA sequence (16p11.2)
met proto-oncogene (hepatocyte growth factor receptor)
distal-less homeo box 2
similar to rat nuclear ubiquituous casein kinase 2
dihydropyrimidinase-like 4
hypothetical protein FLJ20033
inostol 1,4,5-trisphosphate 3-kinase C
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4
somatostatin receptor-interacting protein
brain specific protein
KIAA0769 gene product
PR domain containing 11
hypothetical protein FLJ20510
Consensus includes gb:U66046.1 /DEF=Homo sapiens clone 161455 breast expressed mRNA from chromosome X. /FEA=mRNA /DB_XREF=gi:1519271 /UG=Hs.92663 AF298547
Jun dimerization protein p21SNFT
hypothetical protein FLJ21148
transducin beta-like 1 protein
p21(CDKN1A)-activated kinase 4
hypothetical protein FLJ22004
Consensus includes gb:BF508244 /FEA=EST /DB_XREF=gi:11591542 /DB_XREF=est:U1-H-B14-aqa-b-02-O-U1.s1 /CLONE=IMAGE:3089210 /UG=Hs.300832 ESTs. Highly similar to BF508244
discoidin domain receptor family, member 2
Consensus includes gb:AK024525.1 /DEF=Homo sapiens cDNA: FLJ20872 fls. clone ADKA02604. /FEA=mRNA /DB_XREF=gi:10436827 /UG=Hs.306683 Homo sapiens c AK024525
neuritized-like (Drosophila)
trypsin, alpha
trans-golgi network protein 2
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)
immunoglobulin superfamily, member 6

Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF: AF005082 1.40
 suc1-associated neurotrophic factor target (FGFR signalling adaptor) NM_006654 1.40
 U79287 1.40
 prostate tumor over expressed gene 1 1.40
 Consensus includes gb:AW503390 /FEA=EST /DB_XREF=gi:7118730 /DB_XREF=est:UH-F-BNO-ekw-c02-0-UJ1.1 /CLONE=IMAGE:3078507 /UG=Hs.135258 ESTs. Was AW503390 1.40
 Consensus includes gb:AL137435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp781G0824 (from clone DKFZp781G0824). /FEA=mRNA /DB_XREF=gi:8607994 /UG=Hs.27 AL137435 1.40
 tenascin R (restridin, jensin) NM_003285 1.40
 polymerase (RNA) II (DNA directed) polypeptide J (13.3kD) BE676209 1.40
 glycine amidinotransferase (L-arginine:glycine amidinotransferase) X88401 1.40
 KIAA1039 protein AK000478 1.40
 taste receptor, type 2, member 8 NM_023918 1.40
 gelsolin (amyloidosis, Finnish type) BE675337 1.40
 f-box and leucine-rich repeat protein 6 NM_024555 1.40
 armadillo repeat protein ALEX2 NM_014782 1.40
 hypothetical protein FLJ10851 NM_018245 1.40
 Consensus includes gb:AK023309.1 /DEF=Homo sapiens cDNA FLJ13847 fs, clone THYRO1000852, highly similar to Human branched-chain amino acid aminotransferas AK023909 1.40
 hypothetical protein FLJ22843 NM_025184 1.40
 hypothetical protein FLJ10920 NM_018157 1.40
 mitogen-activated protein kinase kinase 5 D64478 1.40
 g20 protein NM_016210 1.40
 ATPase, Class V, type 10C NM_011136 1.40
 Consensus includes gb:A143878 /FEA=EST /DB_XREF=gi:3665688 /DB_XREF=est:q604f11 x1 /CLONE=IMAGE:1738029 /UG=Hs.128291 phosphodiesterase 10A /FL=g A143878 1.40
 type 1 tumor necrosis factor receptor shedding aminopeptidase regulator AB011097 1.40
 dual specificity phosphatase 9 NM_001395 1.40
 hypothetical protein FLJ00002 AK000002 1.40
 hypothetical protein FLJ21870 NM_023016 1.40
 DKFZP434J154 protein NM_015610 1.40
 slit homolog 3 (Drosophila) AB011538 1.40
 frizzled-related protein NM_001463 1.40
 tenascin XB NM_019105 1.40
 dual adaptor of phosphotyrosine and 3-phosphoinositides NM_014395 1.40
 Wiskott-Aldrich syndrome protein interacting protein AW058622 1.40
 SKI-like X15217 1.40
 spermidine/spermine N1-acetyltransferase M55580 1.40
 Machado-Joseph disease (spinocerebellar ataxia 3, autosomal dominant, ataxin 3) NM_004993 1.40
 phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila) AF012074 1.40
 general transcription factor IIIC, polypeptide 1 (alpha subunit, 220kD) U02818 1.40
 chromosome 17 open reading frame 1A U43383 1.40
 transmembrane protease, serine 2 AF270487 1.40
 RuvB-like 1 (E. coli) M34421 1.40
 Consensus includes gb:AL583528 /FEA=EST /DB_XREF=gi:12952578 /DB_XREF=est:AL583528 /CLONE=CS000004YH21 (5 prime) /UG=Hs.42915 ARP2 (actin-related eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) AL583528 1.40
 TUJ1281-TV protein NM_016575 1.40
 secreted protein of unknown function AA679287 1.40
 lymphocyte-specific protein 1 NM_002339 1.40
 FBJ murine osteosarcoma viral oncogene homolog B NM_006732 1.40
 kinesin family member 13B NM_015254 1.40
 leukotriene C4 synthase NM_000897 1.40
 transketolase-like 1 X91817 1.40
 hypothetical protein FLJ12552 AK024318 1.40
 Fc fragment of IgG binding protein NM_003890 1.40
 testis specific protein, Y-linked M84893 1.40
 complement component 1, q subcomponent binding protein M89039 1.40
 retinoic acid induced 2 NM_021765 1.40
 Consensus includes gb:AL110206.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566N2022 (from clone DKFZp566N2022). /FEA=mRNA /DB_XREF=gi:5817125 /UG=Hs.30 AL110206 1.40
 eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD) NM_003758 1.40
 protodactherin 18 BF222893 1.40
 hypothetical protein NM_013386 1.40
 son of sevenless (Drosophila) homolog 2 L20686 1.40
 Consensus includes gb:AW168915 /FEA=EST /DB_XREF=gi:6400440 /DB_XREF=est:j15h10.x1 /CLONE=IMAGE:2657275 /UG=Hs.283946 Homo sapiens prostate-spec AW168915 1.40
 Consensus includes gb:AA156240 /FEA=EST /DB_XREF=gi:1727858 /DB_XREF=est:Z50c12.x1 /CLONE=IMAGE:5053566 /UG=Hs.286660 Homo sapiens cDNA: FLJ2218 AA156240 1.40
 olfactory receptor, family 5, subfamily V member 1 NM_030876 1.40
 hypothetical protein BF055311 1.40
 nucleosomal binding protein 1 NM_030763 1.40

ras-like protein
interleukin 2
sodium channel, voltage-gated, type X, alpha polypeptide
A kinase (PRKA) anchor protein (gravin) 12
tumor necrosis factor receptor superfamily, member 11a, activator of NFKB
actin A receptor, type IB
serum-inducible kinase
KIAA0643 protein
T cell receptor beta variable 17
chromosome 18 open reading frame 5
Consensus includes gb:AK000156.1 /DEF=Homo sapiens cDNA FLJ20149 fls. clone COL08213. /FEA=mRNA /DB_XREF=gi:7020059 /UG=Hs.272193 Homo sapiens cDNA AK000156
H2B histone family, member E
hypothetical protein FLJ20284
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2
gb:NM_014823.1 /DEF=Homo sapiens KIAA0344, mRNA. /FEA=mRNA /GEN=KIAA0344 /PROD=KIAA0344 gene product /DB_XREF=gi:766208 /NM_014823
SH3-domain binding protein 2
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3
gb:NM_018123.1 /DEF=Homo sapiens hypothetical protein FLJ10517 (FLJ10517), mRNA. /FEA=mRNA /GEN=FLJ10517 /PROD=hypothetical protein FLJ10517 /DB_XREF=NM_018123
zinc finger protein 8 (CAMPX1)
formyl peptide receptor-like 2
epithelial protein lost in neoplasm beta
nuclear factor /B
Consensus includes gb:AIJ4-0483 /FEA=EST /DB_XREF=gi:4083689 /DB_XREF=est:qp51902.x1 /CLONE=IMAGE:1926506 /UG=Hs.132807 Homo sapiens (clone 3.8-1) A.A1346483
Consensus includes gb:AF070579.1 /DEF=Homo sapiens clone 24487 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387951 /UG=Hs.263819 Homo sapiens clone 24487 AF070579
chromosome 1 open reading frame 25
mannosidase, alpha, class 1A, member 2
small EDRK-ict factor 18 (centromeric)
Consensus includes gb:BE541042 /FEA=EST /DB_XREF=gi:3765786 /DB_XREF=est:801064390f.1 /CLONE=IMAGE:3450599 /UG=Hs.23240 Homo sapiens cDNA FLJ13
Rc-like, expressed in neurons (Drosophila)
HMP19 protein
hypothetical protein
Consensus includes gb:AK024108.1 /DEF=Homo sapiens cDNA FLJ14046 fls. clone HEMBA1006461. /FEA=mRNA /DB_XREF=gi:10436408 /UG=Hs.142677 Homo sapie
tripartite motif-containing 33
protein kinase C and casein kinase substrate in neurons 3
retinal pigment epithelium-derived rodopsin homolog
carboxylesterase 3 (brain)
TPA regulated locus
Consensus includes gb:AW072461 /FEA=EST /DB_XREF=gi:6027459 /DB_XREF=est:va06c09.x1 /CLONE=IMAGE:2567536 /UG=Hs.169284 Homo sapiens phosphorib
zytin
gap junction protein, beta 3, 31kD (connexin 31)
microchidia homolog (mouse)
spleen focus forming virus (SFFV) proviral integration oncogene sp1
Consensus includes gb:AQ275374 /DEF=Homo sapiens partial (GVH3) gene for immunoglobulin heavy chain V region, case 1, clone 19 /FEA=CDS /DB_XREF=gi:7573008 /AJ275374
tumor endothelial marker 5 precursor
androgen-induced prostate proliferative shutloff associated protein
protein kinase C, gamma
Consensus includes gb:AW972359 /FEA=EST /DB_XREF=gi:8162205 /DB_XREF=est:EST384450 /UG=Hs.293334 ESTs
carbonic anhydrase VII
KIAA0664 protein
guanine nucleotide binding protein (G protein), alpha 11 (Gq class)
caspase recruitment domain protein 10
KIAA1026 protein
Consensus includes gb:AU144277 /FEA=EST /DB_XREF=gi:11005798 /DB_XREF=est:AU144277 /CLONE=HEMBA1001415 /UG=Hs.296525 Homo sapiens cDNA FLJ101
cold shock domain protein A
angiotensin receptor 2
matrix metalloproteinase 24 (membrane-inserted)
aryl hydrocarbon receptor nuclear translocator 2
gb:NM_018828.1 /DEF=Homo sapiens hypothetical protein PRO2472 (PRO2472), mRNA. /FEA=mRNA /GEN=PRO2472 /PROD=hypothetical protein PRO2472 /DB_XREF=NM_018828
hypothetical protein FLJ23239
hypothetical protein FLJ10770
vasoactive intestinal peptide receptor 2
hypothetical protein FLJ21007
paraneoplastic antigen MA2

sialic acid binding Ig-like lectin 6	D88359	1.39
cyatobrevin, alpha	NM_001392	1.39
peroxisomal trans 2-enooyl CoA reductase	NM_018441	1.39
Sjogren syndrome antigen 8 (autoantigen La)	BC532829	1.39
RFamide-related peptide precursor	NM_022150	1.39
dual specificity phosphatase 3 (vaccinia virus phosphatase VH11-related)	BC002882	1.39
nuclear receptor coactivator 1	U18179	1.39
hypothetical protein FLJ20753	NM_017950	1.39
homeo box D13	NM_000523	1.39
Consensus includes gb:BG251521 /FEA=EST /DB_XREF=gi:127181337 /DB_XREF=est:602363985f1 /CLONE=IMAGE:4472160 /UG=Hs.16193 Homo sapiens mRNA; cdi	BC251521	1.39
hypothetical protein FLJ10803	AJ554945	1.39
hypothetical protein FLJ20761	NM_017954	1.39
CGI-12 protein	NM_015942	1.39
spinocerebellar ataxia 8	AF126749	1.39
hypothetical protein FLJ12168	NM_024682	1.39
Consensus includes gb:AK022999.1 /DEF=Homo sapiens cDNA FLJ12937 fls, clone NT2RP2005020. /FEA=mRNA /DB_XREF=gi:10434712 /UG=Hs.14846 Homo sapient	AK022999	1.39
fibroblast growth factor 1 (ecdic)	X59085	1.39
member of MYST family histone acetyl transferases, homolog of Drosophila MOF	AL050395	1.39
protein tyrosine phosphatase, receptor type, G	NM_002841	1.39
gasulation brain homeo box 1	L11239	1.39
glioma-associated oncogene homolog (zinc finger protein)	NM_005269	1.39
cadherin 11, type 2, OB-cadherin (osteoblast)	D21254	1.39
neuronal PAS domain protein 2	U77970	1.39
hypothetical protein FLJ20513	NM_017855	1.39
signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	NM_005643	1.39
Consensus includes gb:BE87151 /FEA=EST /DB_XREF=gi:10033692 /DB_XREF=est:7d30a1.1.x1 /CLONE=IMAGE:3248732 /UG=Hs.91626 Homo sapiens clone B18 un	BE87151	1.39
membrane-bound transcription factor protease, site 1	BE890314	1.39
Consensus includes gb:AL530331 /FEA=EST /DB_XREF=gi:12793824 /DB_XREF=est:AL530331 /CLONE=CS0DD007YMO6 (3 prime) /UG=Hs.111243 MADS box transcri	AL530331	1.39
colon cancer-associated protein Mtc1	NM_013326	1.39
early growth response 4	NM_001965	1.39
PH domain containing protein in retina 1	AF001563	1.39
collagen, type X, alpha 1 (Schmid metaphyseal chondrodysplasia)	A376003	1.39
Consensus includes gb:AK021741.1 /DEF=Homo sapiens cDNA FLJ11679 fls, clone HEWBA1004807. /FEA=mRNA /DB_XREF=gi:10432986 /UG=Hs.287439 Homo sapie	AK021741	1.39
peptidylprolyl isomerase E (cytoplasmic E)	AF104013	1.39
KIAA0560 gene product	AB011132	1.39
calpain 3, (p94)	AF127764	1.39
pyruvate dehydrogenase kinase, isoenzyme 3	NM_005391	1.39
fibronectin leucine rich transmembrane protein 2	NM_013231	1.39
matrix metalloproteinase 16 (membrane-inserted)	AB009303	1.39
hypothetical protein FLJ22596	NM_025086	1.39
myosin VI	AA877789	1.39
musashi homolog 1 (Drosophila)	AC003982	1.39
heat shock transcription factor 1	NM_005526	1.39
CGI-142	AL133102	1.39
hypothetical protein MGC3103	NM_024036	1.39
hemoglobin, alpha 2	T50399	1.39
solute carrier family 12, (potassium/chloride transporter) member 5	AF208159	1.39
gb:AF130093.1 /DEF=Homo sapiens clone FLB9835 PRO2860 mRNA, complete cds. /FEA=mRNA /PROD=PRO2860 /DB_XREF=gi:1493490 /UG=Hs.302153 Homo sap	AF130093	1.39
S100 calcium binding protein A3	NM_002960	1.39
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)	NM_002608	1.39
Consensus includes gb:BF446180 /FEA=EST /DB_XREF=gi:11511318 /DB_XREF=est:7p33e08.x1 /CLONE=IMAGE:3647703 /UG=Hs.41639 programmed cell death 2	BF446180	1.39
KIAA0573 protein	A827677	1.39
hypothetical protein FLJ11756	NM_024608	1.39
myeloid/lymphoid or mixed-lineage leukemia (t(8;21) translocation, Drosophila); translocated to, 4	NM_024608	1.39
pvt-1 (murine) oncogene homolog, MYC activator	AB016898	1.39
four jointed box 1 (Drosophila)	M34428	1.39
calmodulin 3 (phosphorylase kinase, delta)	NM_014344	1.39
fibronectin leucine rich transmembrane protein 1	NM_005184	1.39
keratin associated protein 1.1	AF169675	1.39
matrix metalloproteinase 24 (membrane-inserted)	NM_030987	1.39
protein kinase, cGMP-dependent, type I	AL121753	1.39
AtPase, Class VI, type 11A	D45864	1.39
Consensus includes gb:N54942 /FEA=EST /DB_XREF=gi:1196282 /DB_XREF=est:y38008.s1 /CLONE=IMAGE:2444983 /UG=Hs.276590 ESTs	BF439472	1.39
	N54942	1.39

T-box 4	NM_018488	1.39
KIAA1614 protein	AB048834	1.39
pro-melanin-concentrating hormone-like 1	S64288	1.39
zinc finger protein 161	AU146275	1.39
epidermal growth factor receptor substrate EPS15R	AV710549	1.39
Consensus includes gb:AU145357 /FEA=EST /DB_XREF=est:AU145357 /CLONE=HEMBA1004611 /UG=Hs.296700 Homo sapiens cDNA FLJ12	AU145357	1.39
hypothetical protein FLJ20624	NM_017906	1.39
collagen, type I, alpha 2	AA788711	1.39
RAN binding protein 7	BG291787	1.39
KIAA0754 protein	AW663712	1.39
Consensus includes gb:AL049263.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F133 (from clone DKFZp564F133) /FEA=mRNA /DB_XREF=gi:4500011 /UG=Hs.30821	AL049263	1.39
hypothetical protein FLJ21125	NM_024627	1.39
thymosin, beta 10	AF060913	1.39
lucosyltransferase 4 (alpha 1(3) lucosyltransferase, myeloid-specific)	M58596	1.39
Consensus includes gb:AL050328 /DEF=Human DNA sequence from clone RP11-145L22 on chromosome 6p21.32-22.2. Contains the gene for myelinoligodendrocyte gly	AL050328	1.39
catenin (cadherin-associated protein), alpha pseudogene 1	Z37894	1.39
coagulation factor II (thrombin) receptor-like 1	NM_005242	1.39
DKFZP564J0206 protein	A927000	1.39
glutamate receptor, ionotropic, kainate 3	NM_000831	1.39
monoamine oxidase A	NM_000240	1.39
kalikrein 7 (chymotryptic, stratum corneum)	NM_005046	1.39
KIAA0902 protein	NM_014927	1.39
hepatocyte growth factor (hepatopolein A; scatter factor)	M77227	1.39
GRB2-related adaptor protein	NM_006813	1.39
EGF-containing fibulin-like extracellular matrix protein 2	AB030855	1.39
hypothetical protein	NM_018985	1.39
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 1	NM_000499	1.39
DnaJ (Hsp40) homolog, subfamily B, member 6	AL355379	1.39
Consensus includes gb:BC396614 /FEA=EST /DB_XREF=gi:13290062 /DB_XREF=est:602459468F1 /CLONE=IMAGE:4581928 /UG=Hs.154999 ESTs, Moderately similar	BG396614	1.39
reslin (Reed-Steinberg cell-expressed intermediate filament-associated protein)	NM_002566	1.39
transmembrane protein 2	NM_013390	1.39
glycine-N-acetyltransferase	NM_005838	1.39
protein-kinase, interferon-inducible double stranded RNA dependent inhibitor, repressor of (P58 repressor)	AF081567	1.39
basement membrane-induced gene	NM_004848	1.39
hypothetical protein MGC5395	NM_024060	1.39
coronin, actin binding protein, 2B	BF839649	1.39
acidic epiklymal glycoprotein-like 1	NM_001131	1.39
Consensus includes gb:AI378979 /FEA=EST /DB_XREF=gi:4188832 /DB_XREF=est:tc0d007.x1 /CLONE=IMAGE:2067085 /UG=Hs.292911 ESTs	AI378979	1.39
EphB1	AF037334	1.39
tubulin, gamma 2	NM_016437	1.39
hypothetical protein FLJ12891	NM_024950	1.39
spondin 1, (f-spondin) extracellular matrix protein	A1855290	1.39
Consensus includes gb:Y14488.1 /DEF=Homo sapiens mRNA for putative 14kD protein containing SHMT homology, clone pUS1215. /FEA=mRNA /PROD=14 kDa protein	Y14488	1.39
PTEN induced putative kinase 1	AF316873	1.39
PTEN induced putative kinase 1	BF432478	1.39
SMA3	NM_006780	1.39
acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain	NM_000017	1.39
H2A histone family, member X	BE901081	1.39
calpain 6	AJ000388	1.39
hypothetical protein FLJ11126	BC005162	1.39
special AT-rich sequence binding protein 1 (binds to nuclear matrix/scaffold-associating DNA's)	NM_002971	1.39
6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF-1)	NM_000281	1.39
Consensus includes gb:AL050145.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586C2020 (from clone DKFZp586C2020) /FEA=mRNA /DB_XREF=gi:4894356 /UG=Hs.22	AL050145	1.39
RAB11B, member RAS oncogene family	AL575337	1.39
vaccinia related kinase 1	NM_003384	1.39
glutaminase	AF097493	1.39
dihydropyrimidinase-like 4	NM_008426	1.39
BAI1-associated protein 3	BE328802	1.39
eukaryotic translation initiation factor 2C, 1	NM_012189	1.39
KIAA0632 protein	AK001615	1.39
polyadenylation binding protein-interacting protein 1	AL136920	1.39
testis specific protein, Y-linked	NM_003308	1.39
like-glycosyltransferase	BF057493	1.39

collagen, type IV, alpha 6
Consensus includes gb:AL117571.1 /DEF=Homo sapiens mRNA; cDNA DKFZ056401172 (from clone DKFZ056401172). /FEA=mRNA /DB_XREF=gi:5912123 /UG=Hs.24
1.39

zeta-chain (TCR) associated protein kinase (70 kD)
Consensus includes gb:AA628440 /FEA=EST /DB_XREF=gi:2540827 /DB_XREF=IMAGE:1032795 /UG=Hs.83623 nuclear receptor subfamily 1, gi:AA628440
1.39

phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 duncle homolog, Drosophila)
Consensus includes gb:AL121975 /FEA=Human DNA sequence from clone RP3-422B11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the AL121975
1.39

zinc finger protein
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

plexin A1
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

KIA00871 protein
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

peptidylprolyl isomerase C (cyclophilin C)
Consensus includes gb:AA628440 /FEA=EST /DB_XREF=gi:2540827 /DB_XREF=IMAGE:1032795 /UG=Hs.83623 nuclear receptor subfamily 1, gi:AA628440
1.39

Consensus includes gb:AA628440 /FEA=EST /DB_XREF=gi:2540827 /DB_XREF=IMAGE:1032795 /UG=Hs.83623 nuclear receptor subfamily 1, gi:AA628440
1.39

Consensus includes gb:AL121975 /FEA=Human DNA sequence from clone RP3-422B11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the AL121975
1.39

acid phosphatase 5, tartrate resistant
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

myosin XVA
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

statilin
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

2,3-bisphosphoglycerate mutase
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

tumor protein 63 kDa with strong homology to p53
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

acetyl-Coenzyme A transporter
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

hypothetical protein FLJ14803
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Consensus includes gb:8F940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC
1.39

epilepsy, progressive myoclonus type 2, Lafora disease (laforin)
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

hypothetical protein FLJ13352
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Thy-1 cell surface antigen
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

hypothetical protein FLJ10178
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

regulator of G-protein signalling 12
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

calcium channel, voltage-dependent, alpha 1I subunit
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

heart and neural crest derivatives expressed 2
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

vanilloid receptor subtype 1
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

KIAA1822 protein
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Rac GTPase activating protein 1
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

tripartite motif-containing 31
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

KIAA0685 gene product
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

SH3 domain binding glutamic acid-rich protein
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

guanylate binding protein 1, interferon-inducible, 67kD
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Consensus includes gb:BE877397 /FEA=EST /DB_XREF=gi:10326173 /DB_XREF=IMAGE:3887938 /UG=Hs.240140 Human DNA sequence fr
1.39

hypothetical protein
Consensus includes gb:BE877397 /FEA=EST /DB_XREF=gi:10326173 /DB_XREF=IMAGE:3887938 /UG=Hs.240140 Human DNA sequence fr
1.39

MAC, mothers against decapentaplegic homolog 5 (Drosophila)
Consensus includes gb:BE877397 /FEA=EST /DB_XREF=gi:10326173 /DB_XREF=IMAGE:3887938 /UG=Hs.240140 Human DNA sequence fr
1.39

hematopoietic PBX-interacting protein
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

extracellular matrix protein 2, female organ and adipocyte specific
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

WD repeat domain 7
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

hypothetical protein FLJ21477
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

CGI-14 protein
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

tetratropocope repeat domain 3
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

vitamin D (1,25-dihydroxyvitamin D3) receptor
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

wingless-type MMTV integration site family, member 3
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

laminin, beta 1
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

neurogenin 2
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

collagen, type IV, alpha 5 (Alport syndrome)
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

nucleolar protein 3 (epiostats repressor with CARD domain)
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

CD8 antigen
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

CD33 antigen, zeta polypeptide (TIT3 complex)
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

arginine vasopressin receptor 1B
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

vanilloid receptor-related osmotically activated channel; OTRPC4 protein
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

hypothetical protein MGC4293
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

pre-B-cell leukemia transcription factor 2
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens
1.39

Consensus includes gb:AL583520 /FEA=EST /DB_XREF=gi:12952562 /DB_XREF=IMAGE:3500024YE13 (5 prime) /UG=Hs.182183 Homo sapiens mR
1.39

cell division cycle 27
Consensus includes gb:AL583520 /FEA=EST /DB_XREF=gi:12952562 /DB_XREF=IMAGE:3500024YE13 (5 prime) /UG=Hs.182183 Homo sapiens mR
1.39

transcobalamin II; macrocytic anemia
Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA: FLJ21342 fls. clone COL02873. /FEA=mRNA /DB_XREF=gi:10437428 /UG=Hs.306739 Homo sapiens cC
1.39

Consensus includes gb:AK022383.1 /DEF=Homo sapiens cDNA: FLJ21342 fls. clone COL02873. /FEA=mRNA /DB_XREF=gi:10437428 /UG=Hs.306739 Homo sapiens cC
1.39

Consensus includes gb:U38460 /DEF=Human RNA binding motif (RBM) gene, partial cds /FEA=CDS /DB_XREF=gi:10613995 /UG=Hs.326802 Human RNA binding motif (f U38460
1.38

heterogeneous nuclear ribonucleoprotein H3 (2H9)	AF132362	1.38
translocase of inner mitochondrial membrane 8 homolog A (yeast)	NM_004085	1.38
double homeobox, 3	NM_012148	1.38
annexin A6	NM_001155	1.38
Consensus includes gb:AK021633.1 /DEF=Homo sapiens cDNA FLJ11571 fls, clone HEMBA1003328 /FEA=mRNA /DB_XREF=gi:50432853 /UG=Hs.306608 Homo sapiens	AK021633	1.38
melanocortin 4 receptor	NM_005912	1.38
hypothetical protein FLJ23033	NM_024686	1.38
solute carrier family 25 (mitochondrial carrier; Graves disease autoantigen), member 16	AI827990	1.38
similar to DNA-directed RNA polymerase I (135 kDa)	NM_019014	1.38
X-prolyl aminopeptidase (aminopeptidase P) 1, soluble	AF195530	1.38
syndecan 1	NM_002997	1.38
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1-like	NM_006222	1.38
ret finger protein-like 2	NM_006605	1.38
hypothetical protein PRO130	NM_018513	1.38
glutamate receptor, metabotropic 1	U31216	1.38
hypothetical protein FLJ22087	NM_022070	1.38
estrogen receptor 2 (ER beta)	AB006589	1.38
ATP-binding cassette, sub-family B (MDR/TAP), member 6	NM_005689	1.38
nucleophosmin (nucleolar phosphoprotein B23, numatrin)	AL353560	1.38
protease, serine, 7 (enterokinase)	NM_002772	1.38
hypothetical protein MGC10848	NM_030569	1.38
KIAA0810 protein	BE972774	1.38
small proline-rich protein 1A	NM_005987	1.38
Consensus includes gb:AK026273.1 /DEF=Homo sapiens cDNA: FLJ22620 fls, clone HS05629 /FEA=mRNA /DB_XREF=gi:10439077 /UG=Hs.306834 Homo sapiens cDI	AK026273	1.38
Consensus includes gb:AL049369.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586D0518 (from clone DKFZp586D0518) /FEA=mRNA /DB_XREF=gi:4500180 /UG=Hs.25	AL049369	1.38
chromosome 19 open reading frame 7	AB028987	1.38
lethal (3) malignant brain tumor l(3)mb1 protein (Drosophila) homolog	U89358	1.38
KIAA1052 protein	AI655714	1.38
Consensus includes gb:AL117549.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586H0123 (from clone DKFZp586H0123) /FEA=mRNA /DB_XREF=gi:5912083 /UG=Hs.26	AL117549	1.38
immunoglobulin lambda locus	X79782	1.38
protocadherin 17	NM_014459	1.38
hypothetical protein FLJ21916	NM_023112	1.38
solute carrier family 2 (facilitated glucose transporter), member 3	AI631159	1.38
thioredoxin reductase beta	AB019695	1.38
KIAA0143 protein	AA805651	1.38
dolichyl-phosphate (UDP-N-acetylglucosamine) N-acetylglucosaminylphosphotransferase 1 (GlcNAc-1-P transferase)	BC000325	1.38
Consensus includes gb:AL080107.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586D0522 (from clone DKFZp586D0522) /FEA=mRNA /DB_XREF=gi:5262532 /UG=Hs.16	AL080107	1.38
ADP-ribosylation factor-like 7	AW450383	1.38
hypothetical protein PP3501	NM_021731	1.38
pancreatic lipase-related protein 1	NM_006229	1.38
hypothetical protein PRO2521	NM_018530	1.38
CCR4-NOT transcription complex, subunit 8	NM_004779	1.38
myeloid/lymphoid or mixed-lineage leukemia (rithorax homolog, Drosophila); translocated to, 10	AF060938	1.38
B lymphoid tyrosine kinase	BC004473	1.38
bromodomain-containing 3	NM_007371	1.38
fatty acid hydroxylase	NM_024306	1.38
carbohydrate kinase-like	NM_013276	1.38
M24537B subtilis phoB, phoA genes corresponding to nucleotides 2017-3334 of M24537 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M24537B	1.38
KIAA1751 protein	AB051538	1.38
solute carrier family 21 (prostaglandin transporter), member 2	NM_005630	1.38
N-acetylglucosamine-phosphate mutase	AV727834	1.38
DKFZP5641122 protein	BC008122	1.38
NAD(P)H dehydrogenase, quinone 1	AI039874	1.38
inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	AW157094	1.38
a disintegrin and metalloproteinase domain 12 (meltrin alpha)	NM_003474	1.38
cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila)	NM_001407	1.38
maltase-glucoamylase (alpha-glucoosidase)	NM_004668	1.38
DKFZP4348204 protein	AL080148	1.38
X03453 Bacteriophage P1 cre recombinase protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	X03453	1.38
serine (or cysteine) proteinase inhibitor, clade H (heat shock protein 47), member 1	NM_004353	1.38
tumor up-regulated CARD-containing antagonist of caspase nine	NM_014959	1.38
filamin C, gamma (actin binding protein 280)	NM_001458	1.38
T-box 1	NM_005992	1.38

dickkopf homolog 1 (Xenopus laevis) 1.37
 plakophilin 3 1.37
 ectonucleoside triphosphate diphosphohydrolase 1 1.37
 cholinergic receptor, nicotinic, beta polypeptide 4 1.37
 Consensus includes gb:AL080180.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M091 (from clone DKFZp434M091) /FEA=mRNA /DB_XREF=gi:5282658 /UG=Hs.1822 AL080180 1.37
 transcription factor 4 1.37
 protein kinase C-like 2 1.37
 dysstrobrevin, alpha 1.37
 neurotrophic tyrosine kinase, receptor, type 3 1.37
 KIAA1655 protein 1.37
 prolidase-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase, Ehlers-Danlos syndrome type VI) 1.37
 v-maf musculoaponeurotic fibrosarcoma oncogene homolog G (avian) 1.37
 small nuclear RNA activating complex, polypeptide 2, 45kD 1.37
 growth hormone 1 1.37
 sentrin/SUMO-specific protease 3 1.37
 nerve growth factor, beta polypeptide 1.37
 gb:NM_019841.1 /DEF=Homo sapiens epithelial calcium channel 1 (ECAC1), mRNA, /FEA=mRNA /GEN=ECAC1 /PROD=epithelial calcium channel 1 /DB_XREF=gi:978961 /DB_XREF=gi:978961 1.37
 siruin silent mating type information regulation 2 homolog 3 (S. cerevisiae) 1.37
 hypothetical protein MGC2376 1.37
 BCL2-associated athanogene 4 1.37
 polydenylate binding protein-interacting protein 1 1.37
 amine oxidase, copper containing 3 (vascular adhesion protein 1) 1.37
 cholinergic receptor, nicotinic, beta polypeptide 3 1.37
 potassium inwardly-rectifying channel, subfamily J, member 3 1.37
 collagen, type IX, alpha 3 1.37
 regulator of G-protein signalling 5 1.37
 KIAA0241 protein 1.37
 endoglin (Osler-Rendu-Weber syndrome 1) 1.37
 transcription factor AP-4 (activating enhancer binding protein 4) 1.37
 adenylylase cyclase 1 (brain) 1.37
 neurotrophic tyrosine kinase, receptor, type 2 1.37
 phosphatidylinositol glycan, class C 1.37
 Consensus includes gb:AA053367 /FEA=EST /DB_XREF=gi:1544893 /DB_XREF=est:75h04.s1 /CLONE=IMAGE:364855 /UG=Hs.51515 Homo sapiens mRNA; cDNA DI AA053367 1.37
 cadherin 18, type 2 1.37
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34 1.37
 mesenchymal stem cell protein DSC43 1.37
 protein phosphatase 1, regulatory (inhibitor) subunit 12A 1.37
 hypothetical protein 1.37
 RAB28, member RAS oncogene family 1.37
 Consensus includes gb:AU158566 /FEA=EST /DB_XREF=gi:5596087 /DB_XREF=est:wm91f10.x1 /CLONE=IMAGE:2443339 /UG=Hs.119683 ESTs 1.37
 chemokine (C-X-C) receptor 1 1.37
 hypothetical protein FLJ12810 1.37
 Consensus includes gb:AL134453 /FEA=EST /DB_XREF=gi:5602840 /DB_XREF=est:DKFZp547J135_s1 /CLONE=DKFZp547J135 /UG=Hs.106890 Homo sapiens clone 2 AL134453 1.37
 integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51) 1.37
 ATPase, Ca++ transporting, plasma membrane 2 1.37
 CGI-116 protein 1.37
 Consensus includes gb:AU158566 /FEA=EST /DB_XREF=gi:11020087 /DB_XREF=est:AU158566 /CLONE=PLACE3000322 /UG=Hs.287635 Homo sapiens cDNA FLJ143 AU158566 1.37
 calcium channel, voltage-dependent, L type, alpha 1D subunit 1.37
 aquaporin 4 1.37
 myotubular myopathy 1 1.37
 hypothetical protein FLJ10945 1.37
 chromosome 21 open reading frame 80 1.37
 collagen, type IV, alpha 3 (Goodpasture antigen) 1.37
 G88 protein 1.37
 hypothetical protein DKFZp566J091 1.37
 calcium channel, voltage-dependent, alpha 1H subunit 1.37
 Consensus includes gb:AK024109.1 /DEF=Homo sapiens cDNA FLJ14047 fs, clone HEMBA1006817 /FEA=mRNA /DB_XREF=gi:10436407 /UG=Hs.301186 Homo sapiens cDNA AK024109 1.37
 Consensus includes gb:AK025029.1 /DEF=Homo sapiens cDNA: FLJ21376 fs, clone COL03231 /FEA=mRNA /DB_XREF=gi:10437464 /UG=Hs.308748 Homo sapiens cDNA AK025029 1.37
 HGC8.2 protein 1.37
 immunoglobulin heavy constant mu 1.37
 KIAA0241 protein 1.37
 GATA binding protein 3 1.37

serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 10
5-hydroxytryptamine (serotonin) receptor 3B
gb:U61168.1 /DEF=Human SH3 domain-containing protein SH3P17 /DB_XREF=gi:14:U61168
Consensus includes gb:AK024784.1 /DEF=Homo sapiens cDNA: FLJ21131 fls, clone CAS06355. /FEA=mRNA /DB_XREF=gi:10437168 /UG=Hs.28264 Homo sapiens mRNA AK024784
nerve growth factor receptor (TNFRSF16) associated protein 1
I factor (complement)
suppression of tumorigenicity 7
hypothetical protein FLJ12816
optic atrophy 1 (autosomal dominant)
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs
dual oxidase 1
kinesin family member 5A
soggy-1 gene
KIAA1001 protein
ribosomal protein L5
hippocalcin
TAFB-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD
hypothetical protein PP3395
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
potassium voltage-gated channel, KQT-like subfamily, member 3
Consensus includes gb:X81637.1 /DEF=H.sapiens clathrin light chain b gene. /FEA=mRNA /DB_XREF=gi:963046 /UG=Hs.73919 clathrin, light polypeptide (Lcb)
myobulatin related protein 8
HSPC067 protein
putative metalloproteinase (family M19)
NGF-A binding protein 2 (EGR1 binding protein 2)
CO163 antigen
angiotensin, ribonuclease, RNase A family, 5
fibrinogen, B beta polypeptide
hypothetical protein FLJ20736
dopamine receptor D2
mel proto-oncogene (hepatocyte growth factor receptor)
paired related homeobox protein
gb:BC002828.1 /DEF=Homo sapiens, Similar to fetal hypothetical protein, clone MGC:3843, mRNA, complete cds. /FEA=mRNA /PROD=Similar to fetal hypothetical protein BC002828
small inducible cytokine subfamily C, member 1 (lymphotactin)
fucosyltransferase 5 (alpha (1,3) fucosyltransferase)
lactin 1
transport-secretion protein 2.2
Consensus includes gb:BG236351 /FEA=EST /DB_XREF=gi:12750198 /DB_XREF=est:na128112.1 /CLONE=IMAGE:4142350 /UG=Hs.148161 ESTs
hypothetical protein
Consensus includes gb:BF032808 /FEA=EST /DB_XREF=gi:10740520 /DB_XREF=est:801455829F1 /CLONE=IMAGE:3859303 /UG=Hs.268784 ESTs
macrophage lectin 2 (calcium dependent)
Consensus includes gb:AK00834.1 /DEF=Homo sapiens cDNA FLJ20827 fls, clone ADKA03543. /FEA=mRNA /DB_XREF=gi:7021152 /UG=Hs.306408 Homo sapiens cDNA AK00834
KIAA061 protein
hypothetical protein DKFZp761F2014
chloride channel, calcium activated, family member 2
chromosome 22 open reading frame 1
Consensus includes gb:AK002179.1 /DEF=Homo sapiens cDNA FLJ11317 fls, clone PLACE1010261, moderately similar to SEGREGATION DISTORTER PROTEIN. /FEA AK002179
hypothetical protein FLJ20509
protein kinase C, alpha binding protein
gb:NM_003483.2 /DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein isoform 1-C (HMGIC), mRNA. /FEA=mRNA /GEN=HMGIC /PROD=high-mobility group protein isoform 1-C (HMGIC), mRNA. /FEA=1269
DKFZP564A122 protein
desmocollin 3
VGF nerve growth factor inducible
peptidyl arginine deiminase, type V
testes-specific heterogeneous nuclear ribonucleoprotein G-T
paired box gene 5 (B-cell lineage specific activator protein)
basoon (presynaptic cytomatrix protein)
interleukin 13
serine/threonine kinase 29
Consensus includes gb:AK025029.1 /DEF=Homo sapiens cDNA: FLJ21376 fls, clone COL03231. /FEA=mRNA /DB_XREF=gi:10437464 /UG=Hs.306748 Homo sapiens cDNA AK025029
voltage-gated sodium channel beta-3 subunit (scn3b gene)
hypothetical protein FLJ23497
BIM88 antigen

calmodulin-like 3	M58026	1.37
KIAA0375 gene product	AB002373	1.37
Consensus includes gb:AL080072.1 /DEF=Homo sapiens mRNA; cDNA DKF Zp564M0618 (from clone DKF Zp564M0618) /FEA=mRNA /DB_XREF=gi:5282482 /UG=Hs.21	AL080072	1.37
ectonucleoside triphosphate diphosphohydrolase 5	NM_001249	1.37
hypothetical protein MG22477	BC001434	1.37
S100 calcium binding protein A9 (calgranulin B)	NM_002965	1.37
KIAA1077 protein	BE500877	1.37
run-related transcription factor 1 (acute myeloid leukemia 1; ami1 oncogene)	L34596	1.37
hypothetical protein PRO0800	NM_016592	1.37
solute carrier family 16 (monocarboxylic acid transporters), member 5	AA705628	1.37
Consensus includes gb:Z21987.1 /DEF=H.sapiens partial cDNA for homologue of mPOU homeobox protein. /FEA=mRNA /PROD=mPOU homeobox protein /DB_XREF=gi:Z21987	Z21987	1.37
vanin 3	NM_018399	1.37
rearranged L-myc fusion sequence	NM_012421	1.37
Z'-5'-oligoadenylate synthetase-like	NM_003723	1.37
KIAA0940 protein	NM_014912	1.37
potential tumor suppressor	NM_013437	1.37
plasma glutamate carboxypeptidase	NM_008102	1.37
hypothetical protein FLJ22714	NM_025050	1.37
angiotensin 1	U83508	1.37
leucine-rich and death domain containing	NM_016494	1.37
small inducible cytokine A5 (RANTES)	NM_002985	1.37
hypothetical protein FLJ13511	AK023573	1.37
jagged 1 (Alagille syndrome)	U81276	1.37
Rho GTPase activating protein 5	AY044631	1.37
matrix metalloproteinase 16 (membrane-inserted)	NM_022584	1.37
gb:NM_007116.1 /DEF=Homo sapiens tenascin XA (TNXA). mRNA. /FEA=mRNA /GEN=TNXA /PROD=tenascin XA /DB_XREF=gi:6005907 /UG=Hs.283750	NM_007116	1.37
BH-prolactin (brain-heart)	NM_002589	1.37
paired basic amino acid cleaving system 4	D87894	1.37
RNA binding motif, single stranded interacting protein 3	NM_014483	1.37
Consensus includes gb:BE551219 /FEA=EST /DB_XREF=gi:5792911 /DB_XREF=est:7656011.x1 /CLONE=IMAGE:3232221 /UG=Hs.16762	BE551219	1.36
group XII secreted phospholipase A2	NM_030821	1.36
matrix metalloproteinase-like 1	NM_004142	1.36
nuclear receptor subfamily 2, group E, member 3	NM_016346	1.36
inparite motif-containing 10	AF220123	1.36
DKFZP566D193 protein	BC005398	1.36
cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)	U17074	1.36
phosphodiesterase 11A	NM_016953	1.36
gonadotropin-regulated testicular RNA helicase	NM_013284	1.36
cyclin-dependent kinase-like 2 /CDC2-related kinase	NM_003948	1.36
lucosyltransferase 6 (alpha (1,3) glucosyltransferase)	U27336	1.36
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	AF070670	1.36
transcription factor 3 (E2A immunoglobulin enhancer binding factor E12/E47)	AF153821	1.36
KIAA1117 protein	A497781	1.36
protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform	BF110421	1.36
alcohol dehydrogenase 1B (class II, beta polypeptide)	AC004770	1.36
hypothetical protein FLJ11136	NM_014458	1.36
EGF-like-domain, multiple 5	NM_019079	1.36
chromosome 1 open reading frame 9	U16797	1.36
Kelch motif containing protein	AF009644	1.36
hypothetical protein	AB71234	1.36
Consensus includes gb:AV691491 /FEA=EST /DB_XREF=gi:10293354 /DB_XREF=est:AV691491 /CLONE=GKCEEB10 /UG=Hs.85335	AV691491	1.36
ephrin-B2	U16797	1.36
E2F transcription factor 3	AF00363	1.36
prodynorphin	NM_024411	1.36
hypothetical protein FLJ13241	NM_025088	1.36
Consensus includes gb:BG389073 /FEA=EST /DB_XREF=gi:13282519 /DB_XREF=est:60241488F1 /CLONE=IMAGE:4523201 /UG=Hs.151010	BG389073	1.36
est variant gene 1	AC004857	1.36
Notch homolog 3 (Drosophila)	NM_000435	1.36
cold autoinflammatory syndrome 1	AK027194	1.36
anterior gradient 2 (Xenopus laevis) homolog	AF088867	1.36
paraoxonase 2	AF001602	1.36
heat shock 70kD protein 1A	NM_005345	1.36
tumor necrosis factor (ligand) superfamily, member 10	NM_003810	1.36

fizzled-related protein	U91903	1.36
G protein-coupled receptor 86	NM_023914	1.36
protein disulfide isomerase-related protein	BE910010	1.36
hypothetical protein FLJ23058	NM_024696	1.36
solute carrier family 9 (sodium/hydrogen exchanger), isoform 2	NM_003048	1.36
homolog of yeast ATP12	AW118608	1.36
cytochrome b-5	M22865	1.36
LUC7-like (S. cerevisiae)	NM_018032	1.36
scrapie responsive protein 1	NM_007281	1.36
disordin domain receptor family, member 1	NM_001954	1.36
Bedilius subtilis /REF=L38424 /DEF=B subtilis dapB, joiF, joiG genes corresponding to nucleotides 2055-2578 of L38424 /LEN=1931 (-5, -M, -3 represent transcript regions)	L38424	1.36
hypothetical protein	A1740515	1.36
relaxin 1 (H1)	BC005956	1.36
KIAA0103 gene product	NM_014873	1.36
TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD	AU05317	1.36
prolactin receptor	AF166329	1.36
G antigen 3	NM_001473	1.36
Cas-B-M (murine) ectropic retroviral transforming sequence b	NM_004351	1.36
vacuolar protein sorting 33B (yeast)	A072363	1.36
cyclic nucleotide gated channel alpha 1	NM_000087	1.36
SMT3 suppressor of mlf two 3 homolog 2 (yeast)	AV726646	1.36
caveolin 1, caveolae protein, 22kD	NM_001753	1.36
paraoxonase 2	NM_000305	1.36
fibroblast growth factor 13	NM_004114	1.36
oxytocin receptor	NM_000916	1.36
peroxisome proliferative activated receptor, gamma	NM_015869	1.36
Consensus includes gb:AF052090.1 /DEF=Homo sapiens clone 23950 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360396 /UG=Hs.106620 Homo sapiens clone 23950	AF052090	1.36
arylsulfatase B	NM_000046	1.36
sialyltransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase)	NM_003033	1.36
protein phosphatase 1, regulatory (inhibitor) subunit 3A (glycogen and sarcoplasmic reticulum binding subunit, skeletal muscle)	NM_002711	1.36
aminohexadecadiene dehydrogenase-phosphopantetheinyl transferase	AF302110	1.36
G antigen 4	NM_001474	1.36
Consensus includes gb:AL049233.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564A023 (from clone DKFZp564A023). /FEA=mRNA /DB_XREF=gi:4499987 /UG=Hs.4224	AL049233	1.36
aldo-keto reductase family 1, member D1 (delta 4-3-ketosteroid-5-beta-reductase)	NM_005989	1.36
peroxisome proliferative activated receptor, delta	NM_006238	1.36
coagulation factor XIII, A1 polypeptide	NM_000129	1.36
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1	U73396	1.36
Consensus includes gb:BC002892.1 /DEF=Homo sapiens, clone IMAGE3940152, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE3940152) /DB_XI	BC002892	1.36
likely ortholog of mouse ganglioside-induced differentiation-associated-protein 2	NM_017886	1.36
CD8 antigen, alpha polypeptide (p32)	AW006735	1.36
chromosome 1 open reading frame 15	AF288395	1.36
cell division cycle 2-like 5 (cholinesterase-related cell division controller)	AJ297710	1.36
hypothetical protein MGC5601	NM_025247	1.36
hypothetical protein FLJ22551	NM_025158	1.36
chromosome 5 open reading frame 3	NM_016691	1.36
Kruppel-like factor 4 (gut)	BF514079	1.36
Consensus includes gb:U80771.1 /DEF=Human EST clone 25267 mariner transposon Hsmar1 sequence. /FEA=mRNA /DB_XREF=gi:2231374 /UG=Hs.245410 Human ES	U80771	1.36
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 duncle homolog, Drosophila)	AF012073	1.36
hepatitis delta antigen-interacting protein A	A583181	1.36
tripartite motif-containing 2	BC005016	1.36
likely ortholog of mouse variant polyadenylation protein CSTF-64	A072408	1.36
hypothetical protein FLJ20602	NM_017896	1.36
Consensus includes gb:AK022892.1 /DEF=Homo sapiens cDNA FLJ12830 fls, clone NT2RP2003073. /FEA=mRNA /DB_XREF=gi:10434549 /UG=Hs.123158 Homo sapier	AK022892	1.36
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	D42072	1.36
hypothetical protein FLJ12691	NM_024572	1.36
ATPase, Na+/K+ transporting, alpha 2 (+) polypeptide	AW440492	1.36
Consensus includes gb:AJ243936.1 /DEF=Homo sapiens mRNA for G16 protein (G16 gene located in the class III region of the major histocompatibility complex). /FEA=m	AJ243936	1.36
Consensus includes gb:AL049325.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D036 (from clone DKFZp564D036). /FEA=mRNA /DB_XREF=gi:4500098 /UG=Hs.1124	AL049325	1.36
zinc finger protein 44 (KOX 7)	A1758888	1.36
N amino acid transporter 3	NM_018018	1.36
gap junction protein, alpha 4, 37kD (connexin 37)	M86789	1.36
small inducible cytokine subfamily A (Cys-Cys), member 22	NM_002990	1.36
melatonin receptor 1B	NM_005859	1.36

KIAA0073 protein	AK025679	1.35
DKFZP564A122 protein	AL110243	1.36
gobg1 phosphoprotein 2	NM_016548	1.36
platelet factor 4	NM_002619	1.36
retinoic acid receptor, beta	NM_016152	1.35
limbic system-associated membrane protein	NM_002338	1.35
Hermansky-Pudlak syndrome	NM_000195	1.35
calcium-sensing receptor (hypocalcemic hyperparathyroidism)	D50855	1.35
hypothetical protein FLJ20202	NM_017709	1.35
adenylate cyclase 9	NM_001116	1.35
meningioma expressed antigen 5 (hyaluronidase)	NM_012215	1.35
discs, large (Orsophila) homolog 4	NM_001365	1.35
bromodomain adjacent to zinc finger domain, 2A	BE513659	1.35
sialic acid binding Ig-like lectin 7	AJ130713	1.35
DKFZP566O1646 protein	AF25793	1.35
neurochondrin	AB018740	1.35
rebaptin-5	NM_004703	1.35
hypothetical protein MGC5347	BE60973	1.35
Consensus includes gb:AV720842 /FEA=EST /DB_XREF=gi:10817994 /DB_XREF=est:AV720842 /CLONE=GLCCOH05 /UG=Hs.282435 ESTs	AV720842	1.35
protein kinase, X-linked	NM_005044	1.35
neurexin 1	AB035356	1.35
solute carrier family 2 (facilitated glucose transporter), member 3	AL110298	1.35
chromodomain helicase DNA binding protein 3	NM_001272	1.35
myosin binding protein H	NM_004997	1.35
trihydrocidic repeat containing 9	AK025084	1.35
lipase, hormone-sensitive	NM_005357	1.35
bradykinin receptor B2	NM_000623	1.35
laminin, gamma 1 (formerly LAMB2)	NM_002293	1.35
TBP-interacting protein	AB014567	1.35
tumor necrosis factor (ligand) superfamily, member 11	AF053712	1.35
HOA14	NM_016484	1.35
glycine receptor, alpha 3	NM_006529	1.35
hypothetical protein DKFZP547M136 similar to widely-interspaced zinc finger motifs	W67467	1.35
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7 (18kD, B18)	M33374	1.35
hypothetical protein FLJ20489	AW149896	1.35
cell division cycle 42 (GTP binding protein, 25kD)	BC003682	1.35
lucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific)	AF305083	1.35
peripheral benzodiazepine receptor-associated protein 1	NM_004756	1.35
hypothetical protein FLJ20087	NM_017662	1.35
Consensus includes gb:AV712602 /FEA=EST /DB_XREF=gi:10731908 /DB_XREF=est:AV712602 /CLONE=DCAACD04 /UG=Hs.5957 Homo sapiens clone 24416 mRNA	AV712602	1.35
purinergic receptor P2X-like 1, orphan receptor	AF065385	1.35
ATP-binding cassette, sub-family A (ABC1), member 4	U88667	1.35
regulator of G-protein signalling 1	NM_002822	1.35
muscle disease-related protein	NM_016298	1.35
arylamide deacetylase (esterase)	NM_001086	1.35
epithelial V-like antigen 1	NM_005797	1.35
LIM homeobox transcription factor 1, beta	NM_002316	1.35
v-est murine thymoma viral oncogene homolog 2	NM_001038	1.35
SRY (sex determining region Y)-box 4	M77196	1.35
hypothetical protein PR02325	BG528420	1.35
Consensus includes gb:BG540048 /FEA=EST /DB_XREF=gi:13532281 /DB_XREF=est:602568675F1 /CLONE=IMAGE:4693242 /UG=Hs.293982 ESTs	NM_018523	1.35
xylokinase (H. influenzae) homolog	BG540048	1.35
Consensus includes gb:AL567940 /FEA=EST /DB_XREF=gi:12921802 /DB_XREF=est:AL567940 /CLONE=CS0DF036YK19 (3 prime) /UG=Hs.7967 ESTs	NM_005108	1.35
caspase recruitment domain protein 14	NM_024110	1.35
pulvative ankyrin-repeat containing protein	NM_015623	1.35
CD84 antigen (leukocyte antigen)	AF054818	1.35
crystallin, beta A4	NM_001886	1.35
cholinergic receptor, muscarinic 5	NM_012125	1.35
Consensus includes gb:AF037219.1 /DEF=Homo sapiens PIX1 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4090971 /UG=Hs.301667 Homo sapiens mRNA; cDNA DKF1	AF037219	1.35
FXYD domain-containing ion transport regulator 2	X86400	1.35
KIAA1042 protein	AK000754	1.35
BAL1-associated protein 2	BC002495	1.35

mitogen-activated protein kinase 8	U34822	1.35
carboxypeptidase, vitellogenic-like	NM_031311	1.35
hypothetical protein PRO2730	N82524	1.35
ADP-ribosyltransferase (NAD ⁺ ; poly (ADP-ribose) polymerase)-like 1	NM_006437	1.35
tenascin XB	M25813	1.35
kinasin family member 59	BF223224	1.35
tumor necrosis factor, alpha-induced protein 2	NM_006291	1.35
calcium channel, voltage-dependent, gamma subunit 5	NM_014404	1.35
adrenergic, alpha-1D-, receptor	M76448	1.35
proedrin P factor, complement	NM_002621	1.35
neutrophil cytosolic factor 2 (65kD, chronic granulomatous disease, autosomal 2)	BC001606	1.35
KIAA0756 protein	A821777	1.35
SCAN domain-containing 2	AF244812	1.35
Consensus includes gb:A185160 /FEA=EST /DB_XREF=est:30g10.s1 /CLONE=IMAGE:1740546 /UG=Hs.326248 Homo sapiens cDNA: FLJ220	A185160	1.35
synovial sarcoma, X breakpoint 3	S82471	1.35
zinc finger protein 291	AF118814	1.35
hypothetical protein FLJ20581	NM_017868	1.35
KIAA0508 protein	AB007975	1.35
echinoderm microtubule-associated protein-like	NM_004434	1.35
paired box gene 5 (B-cell lineage specific activator protein)	NM_016734	1.35
mitogen-activated protein kinase kinase 7	AF006889	1.35
thyroglobulin	NM_003235	1.35
TAL1 (SCL) interrupting locus	NM_003035	1.35
chromosome 12 open reading frame 6	NM_020367	1.35
adrenergic, beta, receptor kinase 1	NM_001819	1.35
uroplekin 3	NM_006953	1.35
corticotropin releasing hormone receptor 1	AF180301	1.35
HIV-1 Tat interactive protein 2, 30 kDa	AF092085	1.35
cancer testis antigen	AJ275978	1.35
glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)	XG9936	1.35
hypothetical protein FLJ12881	AL515916	1.35
Consensus includes gb:AL551046 /FEA=EST /DB_XREF=est:AL551046 /CLONE=CS001065YN13 (5 prime) /UG=Hs.332742 Human DNA from cl	AL551046	1.35
gb:AF165312.1 /DEF=Homo sapiens pre-T-cell receptor alpha chain (PTA) mRNA, alternatively spliced, complete cds. /FEA=CD5 /GEN=PTA /PROD=pre-T-cell receptor a	AF165312	1.35
hypothetical protein FLJ21195 similar to protein related to DAC and cerberus	NM_022469	1.35
hypothetical protein FLJ21324	AW449022	1.35
potassium inwardly-rectifying channel subfamily J, member 13	AJ007557	1.35
Consensus includes gb:AF005081.1 /DEF=Homo sapiens skin-specific protein (kp32) mRNA, partial cds. /FEA=mRNA /GEN=XP32 /PROD=skin-specific protein /DB_XREF=	NM_005764	1.35
epithelial protein up-regulated in carcinoma, membrane associated protein 17	M24915	1.35
CD44 antigen (homing function and Indian blood group system)	NM_024812	1.35
hypothetical protein FLJ12015	U07969	1.35
cadherin 17, LI cadherin (liver-intestine)	NM_007327	1.35
glutamate receptor, ionotropic, N-methyl D-aspartate 1	NM_002784	1.35
RuvB-like 1 (E. coli)	NM_005393	1.35
plexin B3	A056359	1.35
microtubule-associated protein tau	NM_000266	1.35
Norrie disease (pseudoglioma)	AL161996	1.35
ATPase, Class VI, type 11A	NM_030971	1.35
similar to rat tricarboxylate carrier-like protein	N85363	1.35
cyclin-dependent kinase inhibitor 1C (p57, Kip2)	NM_016529	1.35
ATPase, aminophospholipid transporter-like, Class I, type BA, member 2	NM_016236	1.35
alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I; glycolicaciduria; serine-pyruvate aminotransferase)	NM_005371	1.35
methyltransferase-like 1	AL554008	1.35
G protein-coupled receptor 58	NM_001174	1.35
Rho GTPase activating protein 6	NM_001450	1.35
four and a half LIM domains 2	NM_017918	1.35
hypothetical protein FLJ20847	NM_014847	1.35
KIAA0144 gene product	NM_002353	1.35
tumor-associated calcium signal transducer 2	AF038181	1.35
Consensus includes gb:AF038181.1 /DEF=Homo sapiens clone 23607 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2795901 /UG=Hs.167625 Homo sapiens clone 23607	AF038181	1.35
triple functional domain (PTPRF interacting)	AA716657	1.35
presenilin 2 (Alzheimer disease 4)	NM_000111	1.35
solite carrier family 28, member 3	NM_015839	1.35
ficolin (collagen/fibrinogen domain-containing lectin) 2 (hucolin)		1.35

centaurin, gamma 2
hypotheical protein
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 8
putative opioid receptor, neuromedin K (neurokinin B) receptor-like
calcium channel, voltage-dependent, L type, alpha 1B subunit
RAR-related orphan receptor B
ankyrin repeat domain 3
cytochrome P450, subfamily IVF, polypeptide 2
Consensus includes gb:AA551075 /FEA=EST /DB_XREF=gi:2321327 /DB_XREF=est:kn74h06.s1 /CLONE=IMAGE:1019291 /UG=Hs.109438 Homo sapiens clone 24775 r
hypotheical protein FLJ20034
myobubulin related protein 1
SPK1 protein
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1B
actinin, alpha 2
KIAA1035 protein
chemokine binding protein 2
hypotheical protein MGC2592
Consensus includes gb:A050036 /FEA=EST /DB_XREF=gi:3299153 /DB_XREF=est:an31a09.x1 /CLONE=IMAGE:1700248 /UG=Hs.190616 ESTs
KIAA0419 gene product
junctional adhesion molecule 2
zinc finger protein 93 homolog (mouse)
bromodomain-containing 2
restin (Reed-Sternberg cell-expressed intermediate filament-associated protein)
Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog
endothelial differentiation, sphingolipid G-protein-coupled receptor, 1
neurobeachin
cardiac ankyrin repeat protein
Consensus includes gb:AW975638 /FEA=EST /DB_XREF=gi:8166855 /DB_XREF=est:EST387747 /UG=Hs.293490 ESTs
KIAA0241 protein
hemochromatosis
Consensus includes gb:AL122056.1 /DEF=Homo sapiens mRNA; cDNA DKF7Zp434P088 (from clone DKF7Zp434P088). /FEA=mRNA /DB_XREF=gi:6093253 /UG=Hs.2745 /
nucleoporin 88kD
transportin-SR
NK-receptor
hypotheical protein FLJ20079
homeo box B9
aurilil 2
Consensus includes gb:BE673723 /FEA=EST /DB_XREF=gi:10034264 /DB_XREF=est:7d79h10.x1 /CLONE=IMAGE:3279235 /UG=Hs.82719 Homo sapiens mRNA; cDNA
glycophorin A (includes MN blood group)
AKAP-binding sperm protein ropporin
regulator of G-protein signalling 3
Yes-associated protein 1, 65 kDa
M-phase phosphoprotein 6
tripartite motif-containing 22
homeo box A11
fucosyltransferase 8 (alpha (1,6) fucosyltransferase)
claudin 4
G protein-coupled receptor 32
leucine rich repeat (in FLII) interacting protein 2
serine (or cysteine) proteinase inhibitor, clade C (antithrombin), member 1
chromosome 21 open reading frame 7
DKFZP434N181 protein
guanylate binding protein 2, interferon-inducible
gamma-aminobutyric acid (GABA) A receptor, pi
pim-2 oncogene
DKFZP588111 protein
natural killer cell receptor, immunoglobulin superfamily member
insulin-like growth factor binding protein 4
Consensus includes gb:A263044 /FEA=EST /DB_XREF=gi:3871247 /DB_XREF=est:qz29e03.x1 /CLONE=IMAGE:2028316 /UG=Hs.13438 Homo sapiens clone 24626 mF
hypotheical protein
vimentin beta (SH3-containing adaptor molecule-1)
myeloperoxidase
Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fls. clone COL09829. /FEA=mRNA /DB_XREF=gi:10437851 /UG=Hs.306804 Homo sapiens cC AK025352

NM_014914 1.35
NM_013310 1.35
NM_007037 1.35
NM_006879 1.35
NM_000718 1.35
NM_006914 1.35
NM_020639 1.35
D26480 1.35
NM_017630 1.35
AF057354 1.35
NM_014069 1.35
NM_004714 1.35
NM_001103 1.35
AB028958 1.35
NM_001296 1.35
AL121891 1.35
A050038 1.35
NM_014711 1.35
NM_021219 1.35
BC002863 1.35
S78771 1.35
M87501 1.35
BC002836 1.35
NM_001400 1.35
NM_015678 1.35
NM_014391 1.35
AW975638 1.35
BE503381 1.35
AF144243 1.35
AL122056 1.35
Y08613 1.35
AK022910 1.35
AF022048 1.35
NM_017656 1.35
X18172 1.35
NM_017503 1.35
NM_017323 1.35
NM_002099 1.35
NM_017578 1.35
NM_021106 1.35
A1745185 1.35
NM_005792 1.35
AA083478 1.35
NM_005523 1.35
NM_004480 1.35
NM_001305 1.35
NM_001506 1.35
NM_017724 1.35
D29832 1.35
NM_020152 1.35
NM_015493 1.35
NM_004120 1.35
NM_014211 1.35
NM_006875 1.35
AL050131 1.35
NM_007053 1.35
NM_001552 1.35
NM_001552 1.35
NM_019062 1.35
NM_005775 1.35
J02894 1.35
AK025352 1.35

diacylglycerol kinase, alpha (80kD)	NM_001345	1.35
CD164 antigen, siatomucin	NM_006016	1.35
KIAA0773 gene product	NM_014690	1.35
organic cationic transporter-like 3	NM_004256	1.35
Consensus includes gb:AV699347 /FEA=EST /DB_XREF=est:AV699347 /CLONE=GKCBPH07 /UG=Hs.83623 nuclear receptor subfamily 1, group A	AV699347	1.35
pregnancy specific beta-1-glycoprotein 9	M94690	1.35
Wilms' tumour 1-associated protein	AL583811	1.35
lymphocyte antigen 117	AF000426	1.35
hypothetical protein FLJ12665	NM_024803	1.34
Consensus includes gb:AU145746 /FEA=EST /DB_XREF=gi:11007267 /DB_XREF=est:AU145746 /CLONE=HEMBA1005679 /UG=Hs.82193 esterase Diformylglutathione t	AU145746	1.34
Consensus includes gb:AL359052.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMAGE 1968422. /FEA=mRNA /PROD=ITGBL1, Integrin beta-like 1 /D	AL359052	1.34
nuclear receptor subfamily 2, group F, member 1	BC004154	1.34
cyclic nucleotide gated channel alpha.3	NM_001298	1.34
SRY (sex determining region Y)-box 2	AU56682	1.34
phosphoryl-CoA hydroxylase interacting protein	NM_014759	1.34
opioid receptor, delta 1	NM_000911	1.34
hypothetical protein DKFZP434N074	AL096732	1.34
sialic acid binding Ig-like lectin 5	NM_003830	1.34
cystein-rich hydrophobic domain 2	NM_012110	1.34
interleukin 9 receptor	Z84723	1.34
discs, large (Drosophila) homolog 5	BC002915	1.34
LBP protein; likely ortholog of mouse CRTR-1	NM_014553	1.34
cytochrome b5 reductase b5R.2	NM_016229	1.34
synapsin III	NM_003490	1.34
protein predicted by clone 23627	NM_013299	1.34
DKFZP434N093 protein	AL080144	1.34
hypothetical protein FLJ10134	NM_018004	1.34
hypothetical protein FLJ20619	NM_017904	1.34
Consensus includes gb:AL079294.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMAGE 362780. /FEA=mRNA /DB_XREF=gi:5102751 /UG=Hs.29952	AL079294	1.34
insulin-like 5	NM_005478	1.34
protein tyrosine kinase 9	NZ5562	1.34
regulatory factor X, 2 (influences HLA class II expression)	NM_000635	1.34
Consensus includes gb:A1344415 /FEA=EST /DB_XREF=gi:4081621 /DB_XREF=est:cp13b08.x1 /CLONE=IMAGE:1917879 /UG=Hs.156082 ESTs	A1344415	1.34
stress-associated endoplasmic reticulum protein 1, ribosome associated membrane protein 4	BG107676	1.34
NAG-5 protein	NM_016446	1.34
dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-related protein 2)	NM_001922	1.34
chemokine (C-C motif) receptor 5	NM_000579	1.34
Consensus includes gb:AU157915 /FEA=EST /DB_XREF=gi:11019436 /DB_XREF=est:AU157915 /CLONE=PLACE1010102 /UG=Hs.5518 Homo sapiens cDNA FLJ11311	AU157915	1.34
isocitrate dehydrogenase 2 (NADP+), mitochondrial	U52144	1.34
paired box gene 8	A1948472	1.34
tripartite motif-containing 34	NM_021616	1.34
hypothetical protein MGC2306	BC002557	1.34
Consensus includes gb:BE674468 /FEA=EST /DB_XREF=gi:10035007 /DB_XREF=est:7e04e01.x1 /CLONE=IMAGE:3281496 /UG=Hs.12514 Homo sapiens clone 23688 r	BE674468	1.34
hypothetical protein PRO2037	NM_018616	1.34
Consensus includes gb:AU144530 /FEA=EST /DB_XREF=gi:11006051 /DB_XREF=est:AU144530 /CLONE=HEMBA1002189 /UG=Hs.296654 Homo sapiens cDNA FLJ11	AU144530	1.34
KIAA1053 protein	AL117523	1.34
myeloid/lymphoid or mixed-lineage leukemia (t(11b22) translocated to, 4	NM_005936	1.34
Consensus includes gb:AL157837 /DEF=Human DNA sequence from clone RP11-487121 on chromosome 1q31.1-31.3 Contains a ribosomal protein L23A pseudogene, EE	AL157837	1.34
catenin (cadherin-associated protein), beta 1 (88kD)	NM_001904	1.34
Consensus includes gb:W88621 /FEA=EST /DB_XREF=gi:1404293 /DB_XREF=est:zn71d12.r1 /CLONE=IMAGE:417527 /UG=Hs.288924 Homo sapiens cDNA FLJ11392	W88621	1.34
retinal short-chain dehydrogenase/reductase retSDR3	NM_016246	1.34
RAB6B, member RAS oncogene family	AW118072	1.34
cathepsin K (lysosomal acidosis)	NM_000396	1.34
hypothetical protein FLJ11292	NM_018382	1.34
phosphodiesterase 8B	AK023913	1.34
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 9	NM_020249	1.34
four and a half LIM domains 1	U29538	1.34
heparan sulfate 6-O-sulfotransferase	BF512553	1.34
KIAA0143 protein	AW470003	1.34
lipidosis	AB014531	1.34
casease 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	AU719655	1.34
leptin receptor gene-related protein	AW026535	1.34
Cluster Ind. A1742455:wg43605.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2367625 /clone_end=3 /gb=A1742455 /gi=5110743 /ug=Hs.28974 /len=489	A1742455	1.34

potassium voltage-gated channel, shaker-related subfamily, beta member 3
Consensus includes gb:AV687010 /FEA=EST /DB_XREF=gi:10288873 /DB_XREF=est:AV687010 /CLONE=GKCAVC04 /UG=Hs.284651 ESTs
lipin 1
nuclear receptor subfamily 4, group A, member 2
nuclear antigen Sp100
UDP-N-acetylglucosamine-1,3-D-mannoside beta-1,4-N-acetylglucosaminyltransferase IV-homolog
hypothetical protein TCBAPO758
wingless-type MMTV integration site family, member 2B
spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration)
Consensus includes gb:AL050378.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586i1420 (from clone DKFZp586i1420), partial cds. /FEA=mRNA, GEN=DKFZp586i1420 /AL050378
GLI-Kruppel family member GLI2
histone acetyltransferase
opioid growth factor receptor
nuclear receptor subfamily 1, group 1, member 2
BTG family, member 3
KIAA1025 protein
retinoblastoma binding protein 4
Consensus includes gb:Z34282.1 /DEF=H. sapiens (MAR11) MUC5AC mRNA for mucin (partial). /FEA=mRNA /GEN=MUC5AC /PROD=mucin /DB_XREF=gi:563384 /UG=Hs.294277 ESTs
KIAA0914 gene product
Consensus includes gb:A078169 /FEA=EST /DB_XREF=gi:3412577 /DB_XREF=est:z30d11.x1 /CLONE=IMAGE:1678853 /UG=Hs.323511 Homo sapiens cDNA: FLJ231
radixin
Consensus includes gb:AF339787.1 /DEF=Homo sapiens clone IMAGE:205688, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507325 /UG=Hs.326722 Homo sapiens cAF339787
poly (ADP-ribose) glycohydrolase
hypothetical protein FLJ23594
Consensus includes gb:BE671816 /FEA=EST /DB_XREF=gi:10032446 /DB_XREF=est:7a47d01.x1 /CLONE=IMAGE:3221837 /UG=Hs.21921 ESTs, Moderately similar to /BE671816
prefoldin 4
nuclear receptor subfamily 2, group C, member 1
Consensus includes gb:AV700891 /FEA=EST /DB_XREF=gi:10302862 /DB_XREF=est:AV700891 /CLONE=GKCBQD03 /UG=Hs.292477 ESTs
KIAA0564 protein
Consensus includes gb:AK025833.1 /DEF=Homo sapiens cDNA: FLJ22180 fls, clone HRC00938. /FEA=mRNA /DB_XREF=gi:10438467 /UG=Hs.287692 Homo sapiens cAF025833
hypothetical protein FLJ13397
calcium binding protein 1 (calbrain)
synuclein, alpha (non A4 component of amyloid precursor)
tumor protein D52-like 1
synuclein, alpha interacting protein (synphilin)
Consensus includes gb:AK022421.1 /DEF=Homo sapiens cDNA FLJ12359 fls, clone MAMMA 1002355. /FEA=mRNA /DB_XREF=gi:10433814 /UG=Hs.298718 Homo sapiens cAK022421
hypothetical protein FLJ22659
hypothetical protein FLJ11029
hypocretin (orexin) neuropeptide precursor
chromosome 12 open reading frame 5
gb:U87229.1 /DEF=Homo sapiens death adaptor molecule RAIDD-2 mRNA, complete cds. /FEA=CDS /PROD=death adaptor molecule RAIDD-2 /DB_XREF=gi:5051867 /UG=Hs.2987229
TATA box binding protein (TBP)-associated factor, RNA polymerase I, C, 110kD
Slit3-interacting protein
Interleukin 15
cophilin VII
fritzzled homolog 10 (Drosophila)
KIAA0856 protein
Rap1 guanine-nucleotide-exchange factor directly activated by cAMP
hypothetical protein FLJ11808
growth hormone releasing hormone
myosin light chain 2a
carbonic anhydrase I
KIAA0365 gene product
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1
KIAA0942 protein
sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)
hypothetical protein FLJ20378
interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD)
mitogen-activated protein kinase 11
Consensus includes gb:AA654415 /FEA=EST /DB_XREF=gi:2590569 /DB_XREF=est:m03g09.s1 /CLONE=IMAGE:1192096 /UG=Hs.322808 ESTs, Moderately similar to A AA654415
guanylate cyclase activator 1B (relina)
protein tyrosine phosphatase, non-receptor type 12
cullin 5

likely ortholog of mouse Arida1 NM_017610 1.34
Consensus includes gb:A033741 /FEA=EST /DB_XREF=gi:4685071 /DB_XREF=est:128a05.x1 /CLONE=IMAGE:2242064 /UG=Hs.270384 ESTs. Moderately similar to S1 A033741 1.34
tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor) J03225 1.34
signal transducer and activator of transcription 2, 113kd NM_005419 1.34
malic enzyme 3, NADP(+)-dependent, mitochondrial NM_006680 1.34
ribosomal protein S6 kinase, 90kd, polypeptide 5 NM_004755 1.34
Consensus includes gb:AV720514 /FEA=EST /DB_XREF=gi:10817666 /DB_XREF=est:AV720514 /CLONE=GLCGS809 /UG=Hs.282721 ESTs. Weakly similar to ALU7_HI AV720514 1.34
Fc fragment of IgA receptor for U56238 1.34
KIAA0876 protein AW237172 1.34
trypsin beta 1 AF206666 1.34
ubiquitin associated protein AL136733 1.34
CCR4-NOT transcription complex, subunit 2 AL123426 1.34
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) AC002544 1.34
visual system homeobox 1 homolog, CHX10-like (zebrafish) NM_014588 1.34
Consensus includes gb:AL110223.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434M063 /FEA=mRNA /DB_XREF=gi:5817158 /UG=Hs.2745 AL110223 1.34
Integrin, alpha 5 (fibronectin receptor, alpha polypeptide) NM_002205 1.34
Consensus includes gb:AL080111.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586G2222 /FEA=mRNA /DB_XREF=gi:5262538 /UG=Hs.24 AL080111 1.34
KIAA0090 protein D42044 1.34
opsin 1 (cone pigments), medium-wave-sensitive (color blindness, deutan) NM_000513 1.34
retinoic acid receptor, alpha BE383139 1.34
p21 (CDKN1A)-activated kinase 3 AW085556 1.34
mannose-binding lectin (protein A) 1, pseudogene 1 AF018382 1.34
hypothetical protein FLJ20276 NM_017738 1.34
hypothetical protein MGC2601 NM_024042 1.34
solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1 NM_004170 1.34
SEK1 protein NM_014068 1.34
Consensus includes gb:R71245 /FEA=EST /DB_XREF=gi:844762 /DB_XREF=est:y54e05.s1 /CLONE=IMAGE:143072 /UG=Hs.174303 ESTs R71245 1.34
mucin 3B AB038783 1.34
lubbilin, beta, 5 AL567012 1.34
hypothetical protein MGC10946 NM_030572 1.34
Consensus includes gb:AL080161.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434N144 /FEA=mRNA /DB_XREF=gi:5262623 /UG=Hs.2104 AL080161 1.34
chromodomain helicase DNA binding protein 4 U86321 1.34
small inducible cytokine subfamily A (Cys-Cys), member 19 NM_005132 1.34
Rac8p, a meiotic recombination and sister chromatid cohesion phosphoprotein of the rad21p family AL134904 1.34
hypothetical protein FLJ20274 U79275 1.34
hypothetical protein M14087 1.34
Consensus includes gb:M14087.1 /DEF=Human HL14 gene encoding beta-galactoside-binding lectin, 3 end, clone 2 /FEA=mRNA /DB_XREF=gi:187113 /UG=Hs.287389 M14087 1.34
lymphotoxin beta receptor (TNFR superfamily, member 3) NM_002342 1.34
pre-B-cell leukemia transcription factor 2 BC003111 1.34
phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila) NM_000923 1.34
plexin C1 NM_005781 1.34
urtensin 2 NM_021995 1.34
mevalonate kinase (mevalonic aciduria) NM_000431 1.34
Consensus includes gb:S67777 /DEF=estrogen receptor (exon 6) human, tamoxifen-resistant breast tumor 17, Genomic Mutant, 187 nt /FEA=CDS_1 /DB_XREF=gi:45938: S67777 1.34
nurt-related transcription factor 1 (acute myeloid leukemia 1; ami1 oncogene) NM_001754 1.34
development and differentiation enhancing factor 1 NM_018482 1.34
melanoma antigen, family B, 4 NM_002367 1.34
glucose phosphate isomerase AF187554 1.34
carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) X18354 1.34
PRO0644 protein NM_014136 1.34
solute carrier family 10 (sodium/bile acid cotransporter family), member 2 NM_000452 1.33
leukocyte membrane antigen AF202314 1.33
cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) AU154504 1.33
hypothetical protein FLJ14082 NM_025024 1.33
sex comb on midleg homolog 1 NM_012236 1.33
hypothetical protein DKFZp761C07121 NM_017594 1.33
CD28 antigen (Tp44) AF222343 1.33
melanoma antigen, family A, 10 BC004105 1.33
Consensus includes gb:AX025276.1 /DEF=Homo sapiens cDNA: FLJ21623 fs, clone COL07915 /FEA=mRNA /DB_XREF=gi:0437759 /UG=Hs.306791 Homo sapiens cD AX025276 1.33
pre-B-cell leukemia transcription factor 2 BE397715 1.33
hypothetical zinc finger protein FLJ14011 NM_022103 1.33
Consensus includes gb:AW298127 /FEA=EST /DB_XREF=gi:6704763 /DB_XREF=est:U1-H-BW0-ajs-e-05-Q-U1.s1 /CLONE=IMAGE:2733008 /UG=Hs.118832 ESTs. Highly AW298127 1.33
nuclear receptor coactivator 6 interacting protein NM_024831 1.33

single-stranded DNA binding protein 2	NM_012446	1.33
hypothetical protein FLJ2282	NM_024792	1.33
paired basic amino acid cleaving system 4	D26514	1.33
melanoma antigen, family C, 1	AL022152	1.33
itchy homolog E3 ubiquitin protein ligase (mouse)	AB058663	1.33
Consensus includes gb:AV741657 /FEA=EST /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1	AV741657	1.33
Consensus includes gb:AK022170.1 /DEF=Homo sapiens cDNA FLJ12103 fls. clone MAMMA1000009 /FEA=mRNA /DB_XREF=gi:10433505 /UG=Hs.250813 Homo sapiens prostate cancer tumor suppressor	AK022170	1.33
Pu1a1	AU158251	1.33
PP1201 protein	NM_022152	1.33
cholinergic receptor, nicotinic, alpha polypeptide 4	X89745	1.33
vesicle-associated membrane protein 4	BC005974	1.33
par-3 partitioning defective 3 homolog (C. elegans)	NM_019819	1.33
KIA0450 gene product	NM_014638	1.33
N-acetylglucosamine-1-phosphotransferase (E2 component of 2-oxo-glutarate complex)	NM_016256	1.33
dihydroisoamide S-succinyltransferase	AC006530	1.33
KIA00907 protein	NM_014949	1.33
mannosyl (alpha-1,3)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A	NM_012214	1.33
MHC class II transactivator	U18288	1.33
N-myc downstream regulated	NM_006096	1.33
sperm specific antigen 2	NM_006751	1.33
KIA00643 protein	AA207013	1.33
CGI-72 protein	NM_016018	1.33
recombinant binding protein suppressor of hairless (Drosophila)-like	NM_014276	1.33
protocadherin beta 8	NM_019120	1.33
DNA segment on chromosome 6(unique) 2654 expressed sequence	NM_012135	1.33
inhibin, beta A (activin A, activin AB alpha polypeptide)	NM_002192	1.33
keratin, hair, acidic, 6	NM_003771	1.33
solute carrier family 18 (vesicular acetylcholine), member 3	NM_003055	1.33
complement-c1q tumor necrosis factor-related protein; likely ortholog of mouse CORS26 (collagenous repeat-containing sequence of 26-kDa protein)	NM_030945	1.33
death-associated protein kinase 2	NM_014328	1.33
basic helix-loop-helix domain containing, class B, 2	NM_003670	1.33
crystallin, mu	NM_001888	1.33
osteomodulin	NM_005014	1.33
CGI-105 protein	AA273370	1.33
28kD interferon responsive protein	NM_022147	1.33
hypothetical protein FLJ13215	NM_025004	1.33
folistatin	NM_006350	1.33
growth differentiation factor 5 (cartilage-derived morphogenetic protein-1)	NM_000557	1.33
bone morphogenetic protein receptor, type II (serine/threonine kinase)	U25110	1.33
fibulin 1	NM_006487	1.33
thymidine kinase 2, mitochondrial	BE895437	1.33
PRO0132 protein	NM_014116	1.33
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1	NM_014116	1.33
similar to rat integral membrane glycoprotein POM121	AF026132	1.33
keratin 6A	AU158490	1.33
E2F-like protein	J00269	1.33
flavin containing monooxygenase 1	NM_016521	1.33
Consensus includes gb:AK024976.1 /DEF=Homo sapiens cDNA: FLJ1323 fls. clone COL02374 /FEA=mRNA /DB_XREF=gi:10437405 /UG=Hs.75914 Homo sapiens cDNA AK024976	NM_002021	1.33
potassium channel, subfamily K, member 3 (TASK-1)	NM_002246	1.33
regulator of G-protein signalling 4	BC000737	1.33
signal transducer and activator of transcription 1, 91kD	NM_007315	1.33
G1 to S phase transition 2	NM_018094	1.33
solute carrier family 17 (sodium phosphate), member 3	NM_006632	1.33
fibromodulin	NM_002023	1.33
dom-3 homolog 2 (C. elegans)	NM_005510	1.33
neurofilament, heavy polypeptide (200kD)	NM_021076	1.33
oxysterol-binding protein-related protein 1	NM_021076	1.33
Consensus includes gb:AK021614.1 /DEF=Homo sapiens cDNA FLJ11552 fls. clone HEMBA1003021 /FEA=mRNA /DB_XREF=gi:10432830 /UG=Hs.306606 Homo sapiens KIA0280 protein	W19893	1.33
calcium/calmodulin-dependent protein kinase kinase 2, beta	AK021614	1.33
retinoic acid receptor, beta	D87470	1.33
	AK024748	1.33
	NM_004194	1.33
	NM_000665	1.33

prosoducin
 KIAA0810 protein
 lipopolysaccharide binding protein
 solute carrier family 2 (facilitated glucose transporter), member 6
 epithelial membrane protein 1
 sodium channel, nonvoltage-gated 1, delta
 cytochrome P450, subfamily 1C (mephenytoin 4-hydroxylase), polypeptide 18
 putative zinc finger protein from EUROIMAGE 566589
 synaptophysin 3
 Consensus includes gb:AW516297 /FEA=EST /DB_XREF=gi:7154379 /DB_XREF=est:xd64d08.x1 /CLONE=IMAGE:2791215 /UG=Hs.323231 Homo sapiens cDNA FLJ111 AW516297
 hypothetical protein
 Consensus includes gb:BC545769 /FEA=EST /DB_XREF=gi:13544434 /DB_XREF=est:502573042F1 /CLONE=IMAGE:4701250 /UG=Hs.9927 Homo sapiens mRNA; cDN BG545769
 cyclin-dependent kinase inhibitor 1C (p57, Kip2)
 PTH-responsive osteosarcoma B1 protein
 purinergic receptor P2Y, G-protein coupled, 1
 facioscapular dysplasia (Aarskog-Scott syndrome)
 transducin-like enhancer of split 1 (Etsp1) homolog, Drosophila
 hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5
 KIAA0551 protein
 protein phosphatase 1, regulatory (inhibitor) subunit 12B
 cytochrome P450, subfamily 1C (mephenytoin 4-hydroxylase), polypeptide 8
 Consensus includes gb:AK023918.1 /DEF=Homo sapiens cDNA FLJ13856 fls, clone THYRO1000988. /FEA=mRNA /DB_XREF=gi:10436003 /UG=Hs.288489 Homo sapiens
 ras homolog gene family, member
 Consensus includes gb:AK024455.1 /DEF=Homo sapiens mRNA for FLJ00047 protein, partial cds. /FEA=mRNA /GEN=FLJ00047 /PROD=FLJ00047 protein /DB_XREF=gi:AK024455
 lipase, gastric
 hemopoietic cell kinase
 Consensus includes gb:AU148086 /FEA=EST /DB_XREF=gi:11009607 /DB_XREF=est:AU148086 /CLONE=MAMMA1002554 /UG=Hs.302691 Homo sapiens cDNA FLJ12 AU148086
 Consensus includes gb:AA707189 /FEA=EST /DB_XREF=gi:2717117 /DB_XREF=est:32g01.a1 /CLONE=IMAGE:452016 /UG=Hs.298998 Homo sapiens, Similar to hypoa AA707189
 hypothetical protein FLJ11294
 cysteinyl leukotriene CysLT2 receptor, cDNA: PSEC0146 from clone PLACE1006979
 chromosome X open reading frame 1
 A TP-binding cassette, sub-family A (ABC1), member 1
 eukaryotic translation initiation factor 2, alpha kinase 3
 PRO0628 protein
 T-cell lymphoma invasion and metastasis 2
 nuclear receptor co-repressor 2
 alpha2,8-sialyltransferase
 suppressor of K+ transport defect 1
 Consensus includes gb:AI186145 /FEA=EST /DB_XREF=gi:3736783 /DB_XREF=est:q6a06.x1 /CLONE=IMAGE:1742002 /UG=Hs.154338 Homo sapiens clone 24583 tr AI186145
 nuclear respiratory factor 1
 KIAA0878 protein
 sprouty homolog 1, antagonist of FGF signaling (Drosophila)
 protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform
 aspartate beta-hydroxylase
 KIAA0638 protein
 F-box and WD-40 domain protein 1B
 regulator of G-protein signalling 6
 Consensus includes gb:AA527433 /FEA=EST /DB_XREF=gi:2269502 /DB_XREF=est:ng41b03.a1 /CLONE=IMAGE:337325 /UG=Hs.14846 Homo sapiens mRNA; cDNA D AA527433
 brefeldin A-inhibited guanine nucleotide-exchange protein 1
 interleukin 12B (natural killer cell stimulatory factor 2, cytotoxic lymphocyte maturation factor 2, p40)
 KIAA0775 gene product
 transforming growth factor, alpha
 stannocalcin 1
 insulin-like growth factor 1 receptor
 lymphocyte antigen 6 complex, locus E
 Consensus includes gb:BF47272 /FEA=EST /DB_XREF=gi:11512410 /DB_XREF=est:7m47a12.x1 /CLONE=IMAGE:3648695 /UG=Hs.288190 ESTs
 platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit (45kD)
 lamin A/C
 angiotensinogen 2
 neurofilament, heavy polypeptide (200kD)
 Consensus includes gb:BG478428 /FEA=EST /DB_XREF=gi:13410807 /DB_XREF=est:602523839F1 /CLONE=IMAGE:4642353 /UG=Hs.326416 Homo sapiens mRNA; cDN BG478428
 laminin, beta 4
 AF028325
 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
 M76742

	NM_017849	1.33
	BC000052	1.33
	snail 1 homolog, zinc finger protein (Drosophila)	1.33
	ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with mental retardation)	1.33
	mammalian inositol hexakisphosphate kinase 2	1.33
	glioma pathogenesis-related protein	1.33
	glutamate-oxaloacetate transaminase (glutamine synthase)	1.33
	serine protease inhibitor, Kazal type 5	1.33
	ATPase, Ca++ transporting, plasma membrane 2	1.33
	platelet/endothelial cell adhesion molecule (CD31 antigen)	1.33
	transient receptor potential channel 6	1.33
	hematopoietically expressed homeobox	1.33
	serine/threonine kinase with Dbl- and pleckstrin homology domains	1.33
	stromal antigen 2	1.33
	transcription factor-like 4	1.33
	zinc finger protein Z21	1.33
	putative dimethyladenosine transferase	1.33
	light junction protein 3 (zona occludens 3)	1.33
	aminolevulinic acid delta-synthase 2 (sideroblastic anemia)	1.33
	granulylin	1.33
	gbp-BC001407.1 IDEF=Homo sapiens, Similar to cytochrome c-like antigen, clone MGC-23660, mRNA, complete cds.	1.33
	IDEF=mRNA IPROD-Similar to cytochrome c-like anti	1.33
	hypothetical protein FLJ10829	1.33
	nuclear protein, alexis-telangiectasia locus	1.33
	KIAA0409 protein	1.33
	hypothetical protein FLJ23476	1.33
	regulator of G-protein signalling 6	1.33
	tastin specific leucine rich repeat protein	1.33
	topoisomerase (DNA) I	1.33
	protease inhibitor 3, skin-derived (SKALP)	1.33
	cadherin 20, type 2	1.33
	putative ATPase	1.33
	hypothetical protein FLJ12057	1.33
	DHHC1 protein	1.33
	hypothetical protein FLJ12884	1.33
	neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	1.33
	POU domain, class B, transcription factor 1	1.33
	a disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin type 1 motif, 8	1.33
	Consensus includes gb/AU14404.1 /FEA=EST /DB_XREF=ggi:1003562 /DB_XREF=est/AU14404.1 /CLONE=HemBA1000726 /UG=Hs.293834 Homo sapiens cDNA FLJ11,	1.33
	dopa decarboxylase [aromatic L-amino acid decarboxylase]	1.33
	hypothetical protein FLJ23059	1.33
	hydroxyacyl-Coenzyme A dehydratase (trifunctional protein), alpha subunit	1.33
	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	1.33
	Consensus includes gb/AF052145.1 /DEF=Homo sapiens clone 24400 mRNA sequence, /FEA=mRNA /DB_XREF=ggi:3350454 /UG=Hs.159528 Homo sapiens clone 24400 AF052145	1.33
	phosphorylase kinase, gamma 1 (muscle)	1.33
	small inducible cytokine A3	1.33
	lactoperoxidase	1.33
	transforming growth factor, beta receptor III (betaglycan, 300kD)	1.33
	elastase 1, pancreatic	1.33
	apolipoprotein (a) related gene C	1.33
	mitochondrial ribosomal protein S11	1.33
	zinc finger protein 268	1.33
	transmembrane 4 superfamily member 1	1.33
	hepatocyte nuclear factor 3, beta	1.33
	PR domain containing 8	1.33
	a disintegrin and metalloproteinase domain 11	1.33
	Consensus includes gb/AW975050 /FEA=EST /DB_XREF=ggi:8166254 /DB_XREF=est:EST387165 /UG=Hs.293834 ESTs, Weakly similar to ALU4_HUMAN ALU SUBFAM1 AW975050	1.33
	tudor domain containing 1	1.33
	interleukin enhancer binding factor 3, 90kD	1.33
	chromosome 1 open reading frame 10	1.33
	KIAA0417 gene product	1.33
	hypothetical protein MGC13138	1.33
	hypothetical protein FLJ21839 similar to 5-azacytidine induced gene 2	1.33
	NM_022461	1.33

F-box and WD-40 domain protein 7 (archipelago homolog, Drosophila)
 proteasome (prosome, macropain) 26S subunit, ATPase, 4
 hypothetical protein LOC37147
 Consensus includes gb:BF000251 /FEA=EST /DB_XREF=gi:10700526 /DB_XREF=est:7h24e02.x1 /CLONE=IMAGE:3316922 /UG=Hs.227835 KIAA1049 protein
 putative transmembrane protein
 S-phase kinase-associated protein 2 (p45)
 Consensus includes gb:AW68655 /FEA=EST /DB_XREF=gi:8159499 /DB_XREF=est:EST381732 /UG=Hs.168386 ESTs
 photobin-like protein MDS019
 interleukin 24
 lysyl oxidase-like 2
 Consensus includes gb:AU159040 /FEA=EST /DB_XREF=gi:11020561 /DB_XREF=est:AU159040 /CLONE=THYRO1000678 /UG=Hs.296745 Homo sapiens cDNA FLJ131
 Consensus includes gb:AW851559 /FEA=EST /DB_XREF=gi:7947076 /DB_XREF=est:IL3-CT0221-040400-107-E09 /UG=Hs.202695 Human soluble CD44 (CD44) mRNA, AW851559
 fibroblast growth factor 5
 hypothetical protein FLJ22344
 acid-inducible phosphoprotein
 hypothetical protein FLJ10147
 integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide)
 KIAA1564 protein
 calbindin 1, (28KD)
 Consensus includes gb:AK024584.1 /DEF=Homo sapiens cDNA: FLJ20931 fls, done ADSE01282. /FEA=mRNA /DB_XREF=gi:10436896 /UG=Hs.203076 Homo sapiens c
 myeloid cell leukemia sequence 1 (BCL2-related)
 Raf2 interacting protein 8
 NG3 protein
 Consensus includes gb:AB21841 /FEA=EST /DB_XREF=gi:5657805 /DB_XREF=est:wp07a09.x1 /CLONE=IMAGE:2464120 /UG=Hs.212670 ESTs
 guanylate cyclase 2f, retinal
 GATA binding protein 6
 hypothetical protein MGCS287
 epidermal growth factor receptor substrate EPS15R
 Consensus includes gb:BF348061 /FEA=EST /DB_XREF=gi:112956556 /DB_XREF=est:602022810F1 /CLONE=IMAGE:4157907 /UG=Hs.167988 neural cell adhesion mole
 KIAA0461 protein
 calinin (cadherin-associated protein), alpha 1 (102KD)
 hypothetical protein FLJ11560
 cullin 3
 BH3 interacting domain death agonist
 papillomavirus regulatory factor PRF-1
 B melanoma antigen
 KIAA0444 protein
 epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)
 cytosolic ovarian carcinoma antigen 1
 1-acylglycerol-3-phosphate O-acyltransferase 3
 RNA binding motif protein 12
 retinoblastoma binding protein 4
 KIAA0015 gene product
 Consensus includes gb:U63332.1 /DEF=Human super cysteine rich protein mRNA, partial cds. /FEA=mRNA /PROD=super cysteine rich protein /DB_XREF=gi:1480862 /UK
 sodium channel, voltage-gated, type II, beta polypeptide
 KIAA1456 protein
 SWI/SNF related, actin dependent regulator of chromatin, subfamily d, member 3
 Consensus includes gb:M31949 /DEF=Human Ig rearranged mu-chain V-region gene, subgroup VH-III, exon 1 and 2 /FEA=CDS /DB_XREF=gi:185254 /UG=Hs.247923 Hs.
 myelin transcription factor 1-like
 retinoblastoma binding protein 9
 mannosidase, alpha, class 2A, member 2
 Consensus includes gb:AK024123.1 /DEF=Homo sapiens cDNA FLJ14061 fls, clone HEMBB 1000749, /FEA=mRNA /DB_XREF=gi:10436427 /UG=Hs.306684 Homo sapie
 dedicator of cyto-kinesis 1
 Consensus includes gb:BC025063 /FEA=EST /DB_XREF=gi:12411278 /DB_XREF=est:602276408F1 /CLONE=IMAGE:4364304 /UG=Hs.44888 ESTs
 NK3 transcription factor homolog A (Drosophila)
 prospero-related homeobox 1
 neuronin 2
 E1B-55kDa-associated protein 5
 KIAA0300 protein
 chromosome 21 open reading frame 80
 protease, serine, 3 (trypsin 3)
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1
 Consensus includes gb:AC004862 /DEF=Homo sapiens PAC clone RP4-697H17 from Tq11.23-q21.1 /FEA=CDS /DB_XREF=gi:3900856 /UG=Hs.247768 Homo sapiens P
 AC004862

proteasome (prosome, macropain) subunit, beta type, 1
 prolactin receptor
 runt-related transcription factor 2
 RUB-4b protein
 leucine zipper, putative tumor suppressor 1
 Consensus includes gb:AL035406 /DEF=Human DNA sequence from clone RP1-233K16 on chromosome 1p36.21-36.33 Contains the gene KIAA0444, a putative chromod
 likely ortholog of mouse synaptic vesicle glycoprotein 2a
 phosphatidylserine synthase 2
 hypothetical protein FLJ22601
 arachidonate 5-lipoxygenase
 NPHSZ gene (podocin)
 dead ringer (Drosophila)-like 2 (bright and dead ringer)
 type 1 protein phosphatase inhibitor
 G protein-coupled receptor 49
 paraneoplastic cancer-testis-brain antigen
 taste receptor, type 2, member 10
 ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)
 hypothetical protein FLJ12765
 killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2
 hypothetical protein FLJ10381
 mannose-6-phosphate receptor (cation dependent)
 hypothetical protein FLJ14351
 TBC1 (tre-2/USP6, BUB2, cdc16) domain family, member 1
 hypothetical protein FLJ20581
 sarcoma antigen
 Consensus includes gb:U81961 /DEF=Homo sapiens amiloride-sensitive epithelial sodium channel alpha subunit alpha ENaC-1 and alpha ENaC-2 (SCNN1A) genes, allert
 solute carrier family 8 (sodium/calcium exchanger), member 1
 brain cell membrane protein 1
 CGI-43 protein
 Consensus includes gb:AA446466 /FEA=EST /DB_XREF=gi:2159131 /DB_XREF=est:zw65b12.s1 /CLONE=IMAGE:781055 /UG=Hs.193783 Human DNA sequence from c
 actin filament associated protein
 v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)
 5-methyltetrahydrofolate-homocysteine methyltransferase reductase
 actinin, alpha 2
 aldehyde dehydrogenase 3 family, member A2
 fatty-acid-Coenzyme A ligase, long-chain 6
 G protein-coupled receptor 21
 Consensus includes gb:AF088790.1 /DEF=Homo sapiens acylase precursor (ACON) mRNA, nuclear gene encoding mitochondrial protein, partial cds. /FEA=mRNA /GEI
 Consensus includes gb:AL049767 /DEF=Human DNA sequence from clone 172H20 on chromosome 20q12-13.12 Contains genes for serpinogelin I and II and PI3 (proteas
 Consensus includes gb:Z83929 /DEF=Human DNA sequence from clone 272E8 on chromosome Xp22.13-22.31. Contains a pseudogene similar to MDN2-Like P53-binding
 keratin, hair, basic, 3
 hepatocyte nuclear factor 4, alpha
 nuclear protein, marker for differentiated aortic smooth muscle and down-regulated with vascular injury
 phospholipase A2, group VI (cytosolic, calcium-independent)
 neuropilin 1
 hypothetical protein FLJ14031
 HT017 protein
 mitochondrial ribosomal protein L9
 chloride channel 7
 ATP-binding cassette, sub-family G (WHITE), member 4
 similar to rat myomegalin
 flavin containing monooxygenase 4
 integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)
 NADPH-dependent FMN and FAD containing oxidoreductase
 cadherin 11, type 2, OB-cadherin (osteoblast)
 nuclear receptor subfamily 4, group A, member 3
 vitamin D (1,25-dihydroxyvitamin D3) receptor
 AP5-like 1
 hypothetical protein FLJ11259
 SH3-domain GRB2-like 3
 hypothetical protein FLJ10244
 interleukin 3 receptor, alpha (low affinity)
 postmeiotic segregation increased 2-like 4

a disintegrin and metalloproteinase domain 12 (meltrin alpha)
 U1-snRNP binding protein homolog (70kD)
 translocase of inner mitochondrial membrane 17 homolog A (yeast)
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13
 synuclein, alpha (non A4 component of amyloid precursor)
 tomoyin
 Interferon-induced protein 35
 Immunoglobulin heavy constant mu
 PRO1768 protein
 potassium inwardly-rectifying channel, subfamily J, member 2
 mitogen-activated protein kinase 8 interacting protein 1
 Consensus includes gb:AL109809 /DEF=Human DNA sequence from clone RP4-873D20 on chromosome 20. Contains a PTPNS (protein tyrosine phosphatase, non-recap AL109809)
 Consensus includes gb:BE748563 /FEA=EST /DB_XREF=gi:10162555 /DB_XREF=est:60157152111 /CLONE=IMAGE:3838541 /UG=Hs.13413 Homo sapiens clone 2446: BE748563
 makorin, ring finger protein, 2
 cytochrome P450, subfamily IIE (ethanol-inducible)
 collagen, type IV, alpha 3 (Goodpasture antigen)
 hypothetical protein FLJ11383
 proteasome (prosome, macropain) 26S subunit, ATPase, 2
 Consensus includes gb:AF070566.1 /DEF=Homo sapiens clones 24611 and 24705 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387934 /UG=Hs.302042 Homo sapiens
 N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB)
 postmeiotic segregation increased 2-like 3
 Dombrock blood group
 dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)
 integrin, beta 5
 Consensus includes gb:AF003737.1 /DEF=Homo sapiens chromosome 21q22.1 anonymous mRNA sequence. /FEA=mRNA /DB_XREF=gi:2197088 /UG=Hs.302179 Homi AF003737
 gb:AF130057.1 /DEF=Homo sapiens clone FL85539 PRO1454 mRNA, complete cds. /FEA=mRNA /PROD=PRO1454 /DB_XREF=gi:11493420 /UG=Hs.323288 Homo sap AF130057
 hypothetical protein FLJ20489
 Jun D proto-oncogene
 hypothetical protein FLJ20584
 Consensus includes gb:BE044440 /FEA=EST /DB_XREF=gi:8361493 /DB_XREF=est:h045a07.x1 /CLONE=IMAGE:3040356 /UG=Hs.251278 KIAA1201 protein
 KIAA1024 protein
 Consensus includes gb:BE044440 /FEA=EST /DB_XREF=gi:8361493 /DB_XREF=est:h045a07.x1 /CLONE=IMAGE:3040356 /UG=Hs.251278 KIAA1201 protein
 Immunoglobulin heavy constant gamma 3 (G3m marker)
 apolipoprotein L, 5
 pardin, beta
 sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F
 Consensus includes gb:H95283 /FEA=EST /DB_XREF=gi:1102896 /DB_XREF=est:W20604.s1 /CLONE=IMAGE:234366 /UG=Hs.13431 Homo sapiens clone 24790 mRN: H95283
 bromodomain-containing 2
 small protein effector 1 of Cdc42
 Interleukin 17E
 transmembrane, prostate androgen induced RNA
 adenylate kinase 1
 Consensus includes gb:AW405975 /FEA=EST /DB_XREF=gi:6925032 /DB_XREF=est:U1-HF-BL0-acc-a-05-0-U1.r1 /CLONE=IMAGE:3080366 /UG=Hs.247721 Homo saple AW405975
 podocalyxin-like
 glutathione peroxidase 3 (plasma)
 hypothetical protein FLJ20668
 adaptor-related protein complex 1, mu 2 subunit
 adenosine A1 receptor
 bladder cancer overexpressed protein
 ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) 9kD
 hemochromatosis
 hypothetical protein FLJ11220
 taste receptor, type 2, member 13
 DEAD/H (Asp-Glu-Als-Asp/His) box polypeptide 31
 endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 4
 melanoma antigen, family A, 6
 mas-21-like 2 (C. elegans)
 replication protein A complex 34 kd subunit homolog Rpa4
 Inositol 1,4,5-trisphosphate receptor, type 1
 hypothetical protein SBB187
 protein phosphatase 1, regulatory (inhibitor) subunit 12B
 hypothetical protein FLJ20701
 Bacillus subtilis /REF=X04603 /DEF=B subtilis thrC, thrB genes corresponding to nucleotides 288-932 of X04603 /LEN=2073 (-5, -M, -3 represent transcript regions 5 prime X04603
 guanine nucleotide binding protein 11

gb:AF343666.1 /DEF=Homo sapiens translocation associated fusion protein IRTA1IGA1 (IRTA1IGA1) mRNA, complete cds. /FEA=mRNA /GEN=IRTA1IGA1 /PROD=tr AF343666
 uridine monophosphate synthetase (uridine phosphorylase) transferase and oridine-5'-decarboxylase)
 actin like protein
 olfactory receptor, family 7, subfamily C, member 1
 dipeptidylpeptidase VI
 keratin 4
 Consensus includes gb:AK023345.1 /DEF=Homo sapiens cDNA FLJ13283 fs, clone OVARC1001113, highly similar to Homo sapiens diaphanous 1 (HDIA1) mRNA. /FEA=AK023345
 alkaline phosphatase, placental (Regan isozyme)
 hypothetical protein FLJ21522
 phosphoglucomutase 5
 Consensus includes gb:AK025026.1 /DEF=Homo sapiens cDNA: FLJ21373 fs, clone COL03214. /FEA=mRNA /DB_XREF=gi:10437461 /UG=Hs.306747 Homo sapiens cDNA: AK025026
 protein phosphatase 1, regulatory (inhibitor) subunit 14D
 prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase)
 BTG3 associated nuclear protein
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3
 hypothetical protein FLJ22814
 4-hydroxyphenylpyruvate dioxygenase
 synapsin II
 NK3 transcription factor homolog A (Drosophila)
 PCTAIRE protein kinase 3
 POU domain, class 4, transcription factor 2
 DNA fragmentation factor, 45 kD, alpha polypeptide
 a disintegrin and metalloproteinase domain 20
 polycythemia rubra vera 1; cell surface receptor
 hypothetical protein PRO1817
 Huntingtin interacting protein B
 sphingomyelin phosphodiesterase 2, neutral membrane (neutral sphingomyelinase)
 latrophilin
 killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 2
 CUG triplet repeat, RNA binding protein 2
 nitric oxide synthase 3 (endothelial cell)
 DKFZP564B0769 protein
 egi nine homolog 3 (C. elegans)
 solute carrier family 14 (urea transporter), member 2
 transferrin
 hypothetical protein FLJ10375
 Consensus includes gb:AK026155.1 /DEF=Homo sapiens cDNA: FLJ22502 fs, clone HRC11383. /FEA=mRNA /DB_XREF=gi:10438915 /UG=Hs.298140 Homo sapiens cDNA: AK026155
 acyl-Coenzyme A dehydrogenase, long chain
 plasminogen activator, urokinase receptor
 ATP-binding cassette, sub-family C (CFTR/MRP), member 6
 Consensus includes gb:AL162044.1 /DEF=Homo sapiens mRNA: cDNA DKFZp761L0812 (from clone DKFZp761L0812); partial cds. /FEA=mRNA /GEN=DKFZp761L0812 AL162044
 Consensus includes gb:AK025564.1 /DEF=Homo sapiens cDNA: FLJ21911 fs, clone HEP03855. /FEA=mRNA /DB_XREF=gi:10438116 /UG=Hs.306816 Homo sapiens cDNA: AK025564
 KIAA0551 protein
 Consensus includes gb:AF007137.1 /DEF=Homo sapiens clone 23618 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852612 /UG=Hs.302185 Homo sapiens clone 23618 AF007137
 pregnancy-associated plasma protein A
 DKFZP586P0123 protein
 pc32 related 1
 capicua homolog (Drosophila)
 HT021
 Huntingtin interacting protein M
 G-protein coupled receptor
 potassium voltage-gated channel, shaker-related subfamily, member 2
 Werner helicase interacting protein
 crystallin, gamma A
 leptin receptor
 geranylgeranyl diphosphate synthase 1
 Rls
 HEAT-like repeat-containing protein
 crystallin, beta B1
 engrailed homolog 1
 ADP-ribosylator factor-like 5
 Consensus includes gb:BF222018 /FEA=EST /DB_XREF=gi:111129195 /DB_XREF=est:7p41a04.x1 /CLONE=IMAGE:364827 /UG=Hs.289803 ESTs. Moderately similar to BF222018

a disintegrin and metalloproteinase domain 22	AF155381	1.31
carbohydrate (chondroin 6/keratan) sulfotransferase 3	AB017915	1.31
sepin 6	AK026589	1.31
microfibrillar-associated protein 3	R38475	1.31
solute carrier family 25 (mitochondrial carrier, ornithine transporter) member 15	NM_014252	1.31
UDP-Gal4:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4	BF223021	1.31
KIAA0716 gene product	NM_014705	1.31
phosphatidic acid phosphatase type 2A	AB000888	1.31
integrin, beta 5	NM_002213	1.31
Consensus includes gb-AK439416 /FEA=EST /DB_XREF=gi:4304169 /DB_XREF=est:q04b04.x1 /CLONE=IMAGE:2140495 /UG=Hs.165728 ESTs	AI439416	1.31
Rho GTPase activating protein 6	NM_006125	1.31
alkaline phosphatase, liver/bone/kidney	X14174	1.31
discolidin domain receptor family, member 1	L11315	1.31
NPC1 (Niemann-Pick disease, type C1, gene)-like 1	NM_013389	1.31
integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)	U95204	1.31
solute carrier family 4, sodium bicarbonate cotransporter, member 4	NM_003759	1.31
E3 ubiquitin ligase SMURF1	AF199384	1.31
catalase	NM_001752	1.31
collapsin response mediator protein 1	NM_001313	1.31
arginase, type II	NM_001172	1.31
SMAD in the antisense orientation	NM_022001	1.31
Consensus includes gb-AV851559 /FEA=EST /DB_XREF=gi:7947076 /DB_XREF=est:IL3-CT021-040400-107-E09 /UG=Hs.202695 Human soluble CD44 (CD44) mRNA, AW851559	AW851559	1.31
Consensus includes gb-A1810380 /FEA=EST /DB_XREF=gi:5386948 /DB_XREF=est:w87h10.x1 /CLONE=IMAGE:2312707 /UG=Hs.47274 Homo sapiens mRNA, cDNA C A1810380	C A1810380	1.31
Consensus includes gb-BE464789 /FEA=EST /DB_XREF=gi:9510493 /DB_XREF=est:h37a06.x1 /CLONE=IMAGE:3144178 /UG=Hs.27342 ESTs	BE464789	1.31
Consensus includes gb-AL109728.1 /DEF=Homo sapiens mRNA (full length Insert cDNA clone EUROIMAGE 124430, /FEA=EST /DB_XREF=gi:5689792 /UG=Hs.187513 AL109726	AL109726	1.31
hypothetical protein similar to CG7843	AL138752	1.31
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	D50402	1.31
interleukin 1 receptor, type II	NM_004633	1.31
calpastatin	D16217	1.31
FKBP-associated protein	NM_007070	1.31
midline 1 (Opitz/BBB syndrome)	AF041209	1.31
Consensus includes gb-H49077 /FEA=EST /DB_XREF=gi:988918 /DB_XREF=est:y18a05.s1 /CLONE=IMAGE:274180 /UG=Hs.159179 Homo sapiens mRNA sequence, I H49077	I H49077	1.31
hypothetical protein FLJ12681	A003763	1.31
adenylyate kinase 3	NM_013410	1.31
diaca, large (Drosophila) homolog 3 (neuroendocrine-dlg)	NM_021120	1.31
popery protein 3	NM_022361	1.31
gb-BC002792.1 /DEF=Homo sapiens, clone MGC:3963, mRNA, complete cds. /FEA=EST /DB_XREF=gi:128033890 /UG=Hs.81 BC002792	BC002792	1.31
S164 protein	AV757384	1.31
neutrophilic tyrosine kinase, receptor, type 3	S76475	1.31
homeo box D1	NM_024501	1.31
KIAA1888 protein	AL136572	1.31
Consensus includes gb-AK026847.1 /DEF=Homo sapiens cDNA: FLJ23194 fs, clone REC00490, /FEA=EST /DB_XREF=gi:10439802 /UG=Hs.306887 Homo sapiens cDNA, AK026847	AK026847	1.31
chorionic somatomammotropin hormone 1 (placental lactogen)	NM_022640	1.31
activated leucocyte cell adhesion molecule	AA158721	1.31
ectodermal dysplasia 1, anhidrotic	AF061193	1.31
leucy-RNA synthetase, mitochondrial	NM_015340	1.31
aldehyde dehydrogenase 3 family, member A1	NM_000691	1.31
acetylserotonin O-methyltransferase	BC001620	1.31
transcriptional activator of the c-fos promoter	NM_006365	1.31
Spt transcription factor	NM_003112	1.31
chromosome 4 open reading frame 5	BF939178	1.31
eyes absent homolog 2 (Drosophila)	U71207	1.31
tissue inhibitor of metalloproteinase 4	NM_003256	1.31
zinc finger protein 331: zinc finger protein 463	NM_018555	1.31
testis-specific transcript, Y-linked 2	AF000991	1.31
stem cell growth factor, lymphocyte secreted C-type lectin	D86586	1.31
PRO0246 protein	NM_014123	1.31
CGI-119 protein	NM_016056	1.31
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide, Y chromosome	AF000985	1.31
interferon, alpha 16	NM_002173	1.31
deoxyribonuclease I-like 1	NM_006730	1.31
Consensus includes gb-BE738425 /FEA=EST /DB_XREF=gi:10152417 /DB_XREF=est:50157244.T1 /CLONE=IMAGE:3839147 /UG=Hs.241507 ribosomal protein S6	BE738425	1.31
blivertin reductase A	U34877	1.31

sema domain, immunoglobulin domain (Ig), and GPI membrane anchor, (semaphorin) 7A
 SAM domain, SH3 domain end nuclear localisation signals, 1
 GATA binding protein 2
 thyroid hormone receptor, alpha (erythroid leukemia viral (v-erb-a) oncogene homolog, avian)
 proteasome (prosome, macropain) subunit, beta type, 2
 KIAA1199 protein
 basic-helix-loop-helix-PAS protein
 caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)
 leucine-rich, glioma inactivated 1
 glial cells missing homolog a (Drosophila)
 matrix metalloproteinase 20 (matrilysin)
 leucine-rich repeat-containing 3
 tumor necrosis factor receptor superfamily, member 10b
 catenin1 (cadherin-associated protein), alpha 1 (102kD)
 XAGE-1 protein
 hypothetical protein FLJ10922
 cellular retinoic acid binding protein 1
 duodenal cytochrome b
 macrophage scavenger receptor 1
 calyculin-2
 dystrobrevin, beta
 Consensus includes gb:A1808793 /FEA=EST /DB_XREF=gi:5393359 /DB_XREF=est:wt15d05.x1 /CLONE=IMAGE:2350665 /UG=Hs.249239 collagen, type VIII, alpha 2
 fibroblast growth factor 14
 KIAA0530 protein
 hypothetical protein PRO1097
 aquaporin 4
 adipose specific 2
 DKFZP564O243 protein
 N-acetylated alpha-linked acidic dipeptidase 2
 villin 2 (ezrin)
 goliath protein
 calcium binding protein P22
 hypothetical protein FLJ20260
 RAB interacting factor
 Consensus includes gb:S65761.1 /DEF=anti-colorectal carcinoma heavy chain-glycoprotein CANAG-50 specific IgG1 kappa human, 19.9 hybridoma, antibody 1116NS19.9 S65761
 CDK2-associated protein 1
 forkhead box 11
 PRO1892 protein
 multiple PDZ domain protein
 reversal-inducing-cysteine-rich protein with kazal motifs
 myogenin (myogenic factor 4)
 Consensus includes gb:AF090910.1 /DEF=Homo sapiens clone HQ0262. /FEA=mRNA /DB_XREF=gi:6690195 /UG=Hs.236760 Homo sapiens clone HQ0262
 glutamate receptor, ionotropic, kainate 1
 vacuolar protein sorting 11 (yeast)
 vesicle-associated membrane protein 2 (synaptobrevin 2)
 UDP-Gal4betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 3
 Consensus includes gb:AL121890 /DEF=Human DNA sequence from clone RP5-1116H23 on chromosome 20 Contains a novel gene, a 40S ribosomal protein S21 pseudo
 keratin associated protein 9.9
 KIAA1841 protein
 hepatitis A virus cellular receptor 1
 DKFZP568A0522 protein
 Consensus includes gb:AW954107 /FEA=EST /DB_XREF=gi:8143790 /DB_XREF=est:EST366177 /UG=Hs.303394 Homo sapiens cDNA: FLJ22150 fis, clone HRC00109
 MKP-1 like protein tyrosine phosphatase
 lymphoid blast crisis oncogene
 phosphoinositide-3-kinase, catalytic, beta polypeptide
 diuronidase, alpha-L
 sprouty (Drosophila) homolog 4
 casein kinase 1, gamma 2
 Consensus includes gb:AK073448.1 /DEF=Homo sapiens cDNA FLJ13384 fis, clone PLACE1001082, highly similar to Homo sapiens mRNA for lysine-ketoglutarate reducti
 Consensus includes gb:AL122074.1 /DEF=Homo sapiens mRNA, cDNA DKFZP434N0435 (from clone DN7ZP434N0435), partial cds. /FEA=mRNA /GEN=DKFZP434N043
 KIAA1856 protein
 ubiquitin specific protease 9, Y chromosome (Drosophila fat facets related)
 KIAA0453 protein

AF071542 1.31
 NM_022136 1.31
 AL563460 1.31
 M24899 1.31
 BC000268 1.31
 AB033025 1.31
 NM_022123 1.31
 M87507 1.31
 NM_005097 1.31
 AB026493 1.31
 NM_004771 1.31
 NM_030891 1.31
 BC001281 1.31
 D14705 1.31
 NM_020411 1.31
 NM_018273 1.31
 NM_004378 1.31
 NM_024843 1.31
 A1299239 1.31
 NM_022131 1.31
 A1872185 1.31
 A1806793 1.31
 NM_004115 1.31
 BF223237 1.31
 NM_018499 1.31
 D63412 1.31
 NM_006829 1.31
 BC002571 1.31
 AJ012370 1.31
 AF189015 1.31
 AW194816 1.31
 NM_007236 1.31
 NM_017731 1.31
 U74324 1.31
 S65761 1.31
 NM_004642 1.31
 NM_012188 1.31
 AF118086 1.31
 NM_003829 1.31
 AK022887 1.31
 NM_002479 1.31
 AF090910 1.31
 U16125 1.31
 NM_021729 1.31
 NM_014232 1.31
 AB050855 1.31
 AL121890 1.31
 NM_030875 1.31
 AB046881 1.31
 NM_012206 1.31
 AF113007 1.31
 AW954107 1.31
 NM_007026 1.31
 AF126008 1.31
 BC003393 1.31
 A142444.1 1.31
 NM_030964 1.31
 AL530441 1.31
 AK023446 1.31
 AL122074 1.31
 BE552334 1.30
 NM_004654 1.30
 AA524553 1.30

hypothetical protein FLJ10300
 biliverdin reductase A
 Consensus includes gb:AK025026.1 /DEF=Homo sapiens cDNA: FLJ21373 fra, clone COL03214. /FEA=mRNA /DB_XREF=gi:10437461 /UG=Hs.306747 Homo sapiens cDNA
 melatonin receptor 1A
 KIAA1330 protein
 p21(CDKN1A)-activated kinase 6
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 10
 glucagon-like peptide 1 receptor
 carbonyl reductase 3
 nuclear autoantigen
 hypothetical protein PRO1905
 cyclin-dependent kinase 5, regulatory subunit 1 (p35)
 homogentisate 1,2-dioxygenase (homogentisate oxidase)
 KIAA1218 protein
 galanin receptor 1
 opoid receptor, mu 1
 GCIP-interacting protein p29
 Consensus includes gb:AY295387 /FEA=EST /DB_XREF=gi:5702003 /DB_XREF=estUI-H-B/2-shv-e-04-O-UI.s1 /CLONE=IMAGE:2728039 /UG=Hs.196585 ESTs, Wealt
 uromodulin (uronomucoid, Tamm-Horsfall glycoprotein)
 Consensus includes gb:AL049244.1 /DEF=Homo sapiens mRNA: cDNA DKFZ564C163. /FEA=mRNA /DB_XREF=gi:4498979 /UG=Hs.3070 AL049244
 Consensus includes gb:AK025650.1 /DEF=Homo sapiens cDNA: FLJ21997 fra, clone HEP06590. /FEA=mRNA /DB_XREF=gi:10438234 /UG=Hs.306820 Homo sapiens cDNA
 mitochondrial NAD(+)-dependent retinol dehydrogenase 4
 biliverdin reductase A
 H factor 1 (complement)
 PHD finger protein 2
 cyclin-dependent kinase 3
 KIAA0721 protein
 KIAA0605 gene product
 adaptor-related protein complex 1, gamma 1 subunit
 v-kit sarcoma viral oncogene homolog (avian)
 actin filament branching factor 1
 alpha-1-microglobulin/bikunin precursor
 axonal transport of synaptic vesicles
 runt-related transcription factor 1 (ecute myeloid leukemia 1; aml1 oncogene)
 likely ortholog of mouse testis expressed gene 27
 epoxide hydrolase 1, microsomal (xenobiotic)
 hepatocellular carcinoma-associated antigen 127
 zinc finger protein 81 (HFZ20)
 insulin-like growth factor binding protein 3
 KIAA1193 protein
 hypothetical protein FLJ11561
 corticotropin releasing hormone receptor 2
 ADP-ribosylation factor-like 1
 lymphocyte-activation gene 3
 small inducible cytokine subfamily A (Cys-Cys), member 27
 hypothetical protein FLJ21079
 microtubule-associated protein 1A
 hypothetical protein FLJ10188
 ataxia telangiectasia and Rad3 related
 solute carrier family 8 (sodium/calcium exchanger), member 1
 ecdonucleotide pyrophosphatase/phosphodiesterase 3
 homeo box A1
 methionyl aminopeptidase 1
 prostaglandin E receptor 3 (subtype EP3)
 myosin IB
 Consensus includes gb:AU157078 /FEA=EST /DB_XREF=gi:11016599 /DB_XREF=estAU157078 /CLONE=PLACE1006017 /UG=Hs.287580 Homo sapiens cDNA FLJ135
 metastasis-associated 1-like 1
 BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIIB (S.cerevisiae)
 hypothetical protein FLJ13480
 alpha2,3-sialyltransferase
 nuclear receptor subfamily 2, group F, member 6
 hexosaminidase B (beta polypeptide)

G protein-coupled receptor 42
 GLO2 protein
 pre-B-cell leukemia transcription factor 2
 Consensus includes gb:BF056029 /FEA=EST /DB_XREF=est:70703.x1 /CLONE=IMAGE:3443838 /UG=Hs.174910 ESTs. Weakly similar to cyst BF056029
 hypophthalic protein MGC3156
 phospholipid scramblase 1
 frizzled homolog 7 (Drosophila)
 zinc finger protein RINZF
 gap junction protein, beta 3, 31KD (connexin 31)
 myeloid/lymphoid or mixed-lineage leukemia (t(11q24;q24) translocation; Drosophila); translocated to, 4
 lysyl oxidase-like 2
 bone morphogenetic protein 5
 tropomodulin
 death-associated protein kinase 1
 protein kinase H11
 proteasome (prosome, macropain) subunit, alpha type, 7
 SH3-domain binding protein 4
 dynein, cytoplasmic, light intermediate polypeptide 2
 neural cell adhesion molecule 1
 centrosomal protein 2
 Down syndrome critical region gene 4
 neurogenin 3
 Homer, neuronal immediate early gene, 3
 zinc finger protein 202
 taste receptor, type 2, member 1
 a disintegrin and metalloproteinase domain 12 (meltrin alpha)
 hypophthalic protein FLJ11264
 KIAA0820 protein
 UL18 binding protein 2
 hypophthalic protein FLJ11952
 protein tyrosine phosphatase, receptor type, H
 hypophthalic protein FLJ11585
 tritricolide repeat containing 11 (THR-associated protein, 230 kDa subunit)
 double C2-like domains, alpha
 interleukin 1, epsilon
 extracellular link domain-containing 1
 Consensus includes gb:AA631242 /FEA=EST /DB_XREF=est:mq81b04.s1 /CLONE=IMAGE:1158703 /UG=Hs.21349 ESTs. Moderately similar to R AA631242
 uncharacterized gastric protein ZAS5P
 arfcdne homolog 2 (Drosophila)
 Interleukin 1 receptor antagonist
 Rag D protein
 solute carrier family 2 (facilitated glucose transporter), member 3
 homeo box D9
 zinc finger protein 180 (HZF168)
 killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 3
 opsin 3 (enkephalopsin, panopsin)
 Consensus includes gb:BE465829 /FEA=EST /DB_XREF=est:h80n09.x1 /CLONE=IMAGE:3194177 /UG=Hs.150539 Homo sapiens cDNA FLJ137 BE465829
 A kinase (PRKA) anchor protein 7
 uncoupling protein 3 (mitochondrial, proton carrier)
 synapojelin 2
 Consensus includes gb:AF229166.1 /DEF=Homo sapiens docking protein 1-like protein mRNA, partial cds. /FEA=mRNA (PROD=docking protein 1-like protein /DB_XREF=AF229166
 NY-REN-45 antigen
 corticotropin releasing hormone receptor 1
 hypophthalic protein DKFZp434N1923
 UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylglucosaminyltransferase 3 (GanAc-T3)
 olfactory receptor, family 1, subfamily D, member 4
 contactin associated protein-like 2
 kininogen
 Consensus includes gb:AL137624.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M1812 (from clone DKFZp434M1812). /FEA=mRNA /DB_XREF=gi:8808420 /UG=Hs.3C /AL137624
 B9 protein
 small nuclear ribonucleoprotein polypeptide F
 SH3-domain binding protein 1
 nucleosome assembly protein 1-like 1

NM_005305
 NM_020183
 NM_002586
 NM_002586
 NM_024077
 NM_021105
 AI333851
 NM_023929
 AF098730
 AB011358
 AF117949
 NM_021073
 NM_003275
 BC003614
 AF133207
 NM_002792
 AF015043
 AB040861
 M22094
 BC001433
 NM_005867
 NM_020999
 AB071287
 N46430
 NM_016599
 NM_021841
 NM_018371
 AL138712
 NM_025217
 AK022014
 NM_002842
 BF476502
 AI217362
 NM_003586
 NM_014440
 AL574194
 AA631242
 NM_020659
 AV694434
 AW083357
 AL138717
 AA778684
 NM_014213
 NM_013258
 NM_012313
 NM_014322
 NM_014322
 AF161075
 NM_003356
 AK026758
 NM_016121
 AF229166
 X72304
 NM_030974
 BF063271
 NM_003552
 NM_014141
 NM_000893
 NM_000893
 NM_015681
 NM_003095
 NM_018957
 BC002387

pregnancy specific beta-1-glycoprotein 6 M31125 1.30
Consensus includes gb:AW974816 /FEA=EST /DB_XREF=gi:8166019 /DB_XREF=est:EST386021 /UG=Hs.283517 ESTs. Weakly similar to ALU1_HUMAN ALU SUBFAMI1A974816 1.30
KIAA0988 protein AL136608 1.30
tumor necrosis factor (ligand) superfamily, member 13 BF448647 1.30
Consensus includes gb:AK025206.1 /DEF=Homo sapiens cDNA: FLJ21553 fls. clone COL06529 /FEA=mRNA /DB_XREF=gi:10437671 /UG=Hs.306788 Homo sapiens cDNA: AK025206 1.30
hypothetical protein FLJ10819 NM_018156 1.30
Consensus includes gb:AL109708.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMAGE 254679 /FEA=mRNA /DB_XREF=gi:5689825 /UG=Hs.284310 AL109708 1.30
KIAA0903 protein AB020710 1.30
Consensus includes gb:AL137285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D2416 (from clone DKFZp434D2416) /FEA=mRNA /DB_XREF=gi:5607740 /UG=Hs.30 AL137285 1.30
chitinin (chitinase) 2 AK026415 1.30
nitric oxide synthase 1 (neuronal) U31466 1.30
zinc finger protein 20 (K0X13) AU154474 1.30
tenascin XB X71923 1.30
exotoxins (multiple)-like 3 BC006383 1.30
Consensus includes gb:AL080134.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434G043 (from clone DKFZp434G043) /FEA=mRNA /DB_XREF=gi:5262575 /UG=Hs.1687 AL080134 1.30
C-terminal binding protein 2 BC002486 1.30
diubiquitin NM_006398 1.30
EphB3 NM_004443 1.30
solute carrier family 6 (neurotransmitter transporter, creatine), member 10 U41163 1.30
ATP-binding cassette, sub-family G (WHITE), member 1 U34919 1.30
folate receptor 2 (fetal) NM_000803 1.30
HMT1 hnRNP methyltransferase-like 3 (S. cerevisiae) NM_018854 1.30
splicing factor 30, survival of motor neuron-related BF224259 1.30
ectodermal dysplasia 1, anhidrotic NM_001359 1.30
pyruvate kinase, liver and RBC M15465 1.30
estrogen receptor 1 AF238450 1.30
formyl peptide receptor-like 1 M88107 1.30
hypothetical protein FLJ10811 NM_018228 1.30
calpastatin BG111635 1.30
hypothetical protein FLJ13381 NM_025068 1.30
KIAA0310 protein NM_025154 1.30
zinc finger protein 232 NM_014519 1.30
GAS2-related on chromosome 22 Y07846 1.30
mucin 3A, intestinal AF007194 1.30
Consensus includes gb:AK026465.1 /DEF=Homo sapiens cDNA: FLJ22812 fls. clone KIAA2955 /FEA=mRNA /DB_XREF=gi:10439335 /UG=Hs.306853 Homo sapiens cDNA: AK026465 1.30
hydroxysteroid (11-beta) dehydrogenase 2 NM_000196 1.30
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated) X98743 1.30
catenin (cadherin-associated protein), alpha 2 NM_004389 1.30
interferon, alpha 2 M54886 1.30
thionexin-like 2 NM_006541 1.30
amphotropic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 AV705253 1.30
aldehyde dehydrogenase 9 family, member A1 NM_000696 1.30
unc53 (C. elegans) homolog B NM_030930 1.30
valosin-containing protein W60953 1.30
adenosine deaminase, RNA-specific, B1 (homolog of rat RED1) NM_001112 1.30
par-3 partitioning defective 3 homolog (C. elegans) NM_01112 1.30
hypothetical protein LOC57333 AF191685 1.30
chromosome 1 topen reading frame 9 NM_013279 1.30
timeless homolog (Drosophila) AK000721 1.30
RAB1, member RAS oncogene family AL530284 1.30
thyroid stimulating hormone, beta NM_000549 1.30
hypothetical protein MGC11271 NM_024323 1.30
claudin 10 NM_006984 1.30
translocase of inner mitochondrial membrane 17 homolog A (yeast) AK023063 1.30
Consensus includes gb:AK027151.1 /DEF=Homo sapiens cDNA: FLJ23498 fls. clone LNC02683 /FEA=mRNA /DB_XREF=gi:10440206 /UG=Hs.306904 Homo sapiens cDNA: AK027151 1.30
hypothetical protein FLJ22390 NM_022746 1.30
apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1 NM_001644 1.30
transforming growth factor, beta 3 J03241 1.30
Consensus includes gb:AA587390 /FEA=EST /DB_XREF=est:tm82h01.s1 /CLONE=IMAGE:1090417 /UG=Hs.321110 ESTs. Highly similar to mucin 8 1.30
ret finger protein-like 3 antisense AL021837 1.30
hypothetical protein FLJ14001 NM_024677 1.30
B9 protein BC002944 1.30
Consensus includes gb:AK028723.1 /DEF=Homo sapiens cDNA: FLJ23070 fls. clone LNC05829 /FEA=mRNA /DB_XREF=gi:10439641 /UG=Hs.305579 Homo sapiens cDNA: AK028723 1.30

[illegible]

Consensus includes gb:AF149193 /FEA=EST /DB_XREF=gi:127457 /DB_XREF=est:at40e04.x1 /CLONE=IMAGE:2374494 /UG=Hs.17639 Homo sapiens ubiquitin protein AL117566
 ubiquitin-activating enzyme E1C (UBA3 homolog, yeast) BF796940
 Consensus includes gb:BF796940 /FEA=EST /DB_XREF=gi:12101984 /DB_XREF=est:602258153F1 /CLONE=IMAGE:4341588 /UG=Hs.294100 ESTs NM_001868
 carboxypeptidase A1 (pancreatic) AW024233
 glycine-N-acyltransferase U05682
 TYRO3 protein tyrosine kinase U81640
 survival motor neuron pseudogene M30646
 erythrocyte membrane protein band 4.2 AF225881
 ATPase, Ca++ transporting, type 2C, member 1 A4555068
 hypothetical protein FLJ22087 M73554
 cyclin D1 (PRAD1; parathyroid adenomatosis 1) D32039
 chondroilin sulfate proteoglycan 2 (versican) AK026793
 Consensus includes gb:AK026793.1 /DEF=Homo sapiens cDNA: FLJ23140 fs, clone LNG09065, /FEA=mRNA /DB_XREF=gi:10439729 /UG=Hs.287728 Homo sapiens cDNA: AK026793
 Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse) M92424
 potassium voltage-gated channel, shaker-related subfamily, beta member 2 AL520102
 Consensus includes gb:AF12086 /FEA=EST /DB_XREF=est:wd8408.x1 /CLONE=IMAGE:2338311 /UG=Hs.194750 Human DNA sequence from cDNA: AF12086
 syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan) BE048514
 Consensus includes gb:AW301235 /FEA=EST /DB_XREF=gi:8710912 /DB_XREF=est:xs69c06.x1 /CLONE=IMAGE:2774890 /UG=Hs.66185 Homo Sapiens mRNA, partial AW301235
 gqij associated, gamma adaptin ear containing, ARF binding protein 2 AW088154
 CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) L25259
 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit 9 AA917672
 chromosome 11 open reading frame 14 NM_020645
 karyopherin alpha 8 (importin alpha 7) BG403834
 Cluster incl. A1738452; w32a07.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2391924 /clone_end=3 /gb=AF1738452 /gb=5100433 /ug=Hs.203286 /len=533
 leucine rich repeat (in FLII) interacting protein 1 AF130054
 hydroxyteroid (17-beta) dehydrogenase 7 NM_016371
 formyl peptide receptor 1 M60626
 protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform AJ271832
 glycoprotein M8A BF939489
 olfactory receptor, family 7, subfamily E, member 18 pseudogene AC096271
 Consensus includes gb:AF207792 /FEA=EST /DB_XREF=gi:3769734 /DB_XREF=est:so89403.x1 /CLONE=IMAGE:1953029 /UG=Hs.306019 ESTs, Weakly similar to ALU7 NM_022735
 gqij phosphoprotein 1 NM_012189
 fibronectin II NM_006715
 mammoside, alpha, class 2C, member 1 AF041811
 ets variant gene 6 (TEL oncogene) M19720
 v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian) 129
 Consensus includes gb:AF207792 /FEA=EST /DB_XREF=gi:3769734 /DB_XREF=est:so89403.x1 /CLONE=IMAGE:1953029 /UG=Hs.306019 ESTs, Weakly similar to ALU7 NM_022735
 KIAA0873 protein 129
 Consensus includes gb:AK024457.1 /DEF=Homo sapiens mRNA for FLJ00049 protein, partial cds, /FEA=mRNA /GEN=KRTAP2.1A /PROD=keratin associated protein K NM_030977
 Consensus includes gb:AU147820 /FEA=EST /DB_XREF=gi:11009141 /DB_XREF=est:AU147820 /CLONE=MAMMA1001206 /UG=Hs.288490 Homo sapiens cDNA FLJ12 AU147820
 double-stranded RNA specific adenosine deaminase NM_018702
 hypothetical protein FLJ13031 NM_024688
 Consensus includes gb:AU039469 /FEA=EST /DB_XREF=gi:3278663 /DB_XREF=est:ox41a08.s1 /CLONE=IMAGE:1658870 /UG=Hs.27362 ESTs AK039469
 aquaporin 8 NM_001169
 Consensus includes gb:AK024955.1 /DEF=Homo sapiens cDNA: FLJ21342 fs, clone COL02873, /FEA=mRNA /DB_XREF=gi:10437426 /UG=Hs.308739 Homo sapiens cDNA: AK024955
 a disintegrin and metalloproteinase domain 7 AF215824
 phospholipase D1, phosphatidylcholine-specific U38545
 hypothetical protein DKFZp434C0923 NM_017598
 Consensus includes gb:AL035587 /DEF=Human DNA sequence from clone RP3-475N16 on chromosome 6p12.3-21.2. Contains the genes for CTGA4, pre-T cell receptor, BF115148
 progesterin induced protein AF119817
 discs, large (Drosophila) homolog-associated protein 2 NM_024689
 hypothetical protein FLJ14103 AF119817
 Consensus includes gb:AF052146.1 /DEF=Homo sapiens clone 24653 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3380455 /UG=Hs.155085 Homo sapiens clone 24653 AF052146
 KIAA0523 protein BF115148
 disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila) NM_001343
 flavin containing monooxygenase 5 NM_001461
 hepatocyte nuclear factor 3, beta AB028021
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1 NM_003069
 tripartite motif-containing 10 AF220122
 cleavage and polyadenylation specific factor 1, 160KD subunit BE676642
 hypothetical protein DKFZp782B226 AK001821
 galactosylceramidase (Krabbe disease) NM_000153
 chromosome 1 open reading frame 14 NM_030933

protein tyrosine phosphatase, receptor type, F 1.29
 Interleukin 9 receptor 1.29
 death-associated protein kinase 1 1.29
 hypothetical protein PRQ1787 1.29
 signal transducer and activator of transcription 1, 91KD 1.29
 olfactory receptor, family 1, subfamily E, member 2 1.29
 surfactant protein A binding protein 1.29
 cullin 4A 1.29
 KIAA0918 protein 1.29
 Bacteriophage JREF=X03453 /DEF=Bacteriophage P1 cre recombinase protein corresponding to nucleotides 581-1001 of X03453 /LEN=1058 (-5 and -3 represent transcrip 1.29
 Consensus includes gb:AF035314.1 /DEF=Homo sapiens clone 23651 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2661076 /UG=Hs.134526 Homo sapiens clone 23651 AF035314 1.29
 claudin 18 1.29
 DNFP2434C212 protein 1.29
 ephrin-A2 1.29
 chondroitin sulfate proteoglycan BEHAB/brevican 1.29
 KIAA0231 protein 1.29
 aldo-keto reductase family 1, member B10 (aldose reductase) 1.29
 dynein, axonemal, heavy polypeptide 17 1.29
 Consensus includes gb:AF49382 /FEA=EST /DB_XREF=gi:989223 /DB_XREF=est:q19506.s1 /CLONE=IMAGE:274115 /UG=Hs.124984 ESTs, Moderately similar to ALU7. H49382 1.29
 collagen, type XIII, alpha 1 1.29
 Ras suppressor protein 1 1.29
 frizzled homolog 7 (Drosophila) 1.29
 secreted protein, acidic, cysteine-rich (osteonection) 1.29
 kinesin family member 3B 1.29
 sarcosine dehydrogenase 1.29
 hypothetical protein FLJ21687 1.29
 anaplastic lymphoma kinase (K1-1) 1.29
 adaptor protein containing PH domain, PTB domain and leucine zipper motif 1.29
 Consensus includes gb:BF431618 /FEA=EST /DB_XREF=gi:11443732 /DB_XREF=est:7016e03.x1 /CLONE=IMAGE:3574349 /UG=Hs.12342 Homo sapiens clone 24538 r BF431618 1.29
 glutamate receptor, ionotropic, N-methyl D-aspartate 2C 1.29
 Consensus includes gb:AL050335 /DEF=Human DNA sequence from clone RP1-190J20 on chromosome 6p22.3-24.1. Contains ESTs, STSs and GSSs. 1.29
 cytochrome P450, subfamily IIE (ethanol-inducible) 1.29
 edoquinone A3 receptor 1.29
 hypothetical protein FLJ20234 1.29
 alpha-2-glycoprotein 1, zinc 1.29
 Consensus includes gb:AF070577.1 /DEF=Homo sapiens clone 24461 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387949 /UG=Hs.4817 Homo sapiens clone 24461 m AF070577 1.29
 1-acetylglucosyl-3-phosphate O-acetyltransferase 1 (lysophosphatidic acid acyltransferase, alpha) 1.29
 eukaryotic translation initiation factor 2, subunit 1 (alpha, 35KD) 1.29
 giant axonal neuropathy (giganonin) 1.29
 CD209 antigen-like 1.29
 Fc fragment of IgE, low affinity II, receptor for (CD23A) 1.29
 hypothetical protein FLJ21634 1.29
 Consensus includes gb:AL034396 /DEF=Human DNA sequence from clone 1158B12 on chromosome Xp11.21-11.4 Contains the ZXDA gene for X-linked duplicated Zinc fl 1.29
 transforming growth factor, beta 2 1.29
 polycystic kidney disease 1 (autosomal dominant) 1.29
 chromosome 1 open reading frame 8 1.29
 putative integral membrane transporter 1.29
 Consensus includes gb:AK022362.1 /DEF=Homo sapiens cDNA FLJ12300 fls, clone MAMMA1001854. /FEA=mRNA /DB_XREF=gi:10433744 /UG=Hs.287510 Homo sapie 1.29
 tumor-associated calcium signal transducer 1 1.29
 Consensus includes gb:AK024871.1 /DEF=Homo sapiens cDNA: FLJ21218 fls, clone COL00537. /FEA=mRNA /DB_XREF=gi:10437282 /UG=Hs.306712 Homo sapiens c 1.29
 TUGA protein 1.29
 H1 histone family, member 5 1.29
 fatty-acid-Coenzyme A ligase, long-chain 5 1.29
 histamine N-methyltransferase 1.29
 synuclein, gamma (breast cancer-specific protein 1) 1.29
 claudin 6 1.29
 KIAA0877 protein 1.29
 putative DNA/chromatin binding motif 1.29
 Consensus includes gb:AW241752 /FEA=EST /DB_XREF=gi:6575506 /DB_XREF=est:xt74f04.x1 /CLONE=IMAGE:2700223 /UG=Hs.160728 Homo sapiens cDNA FLJ1116 AW241752 1.29
 gap junction protein, beta 5 (connexin 31.1) 1.29
 matrix metalloproteinase 14 (membrane-inserted) 1.29
 hypothetical protein FLJ22021 1.29
 attractin 1.29

plakophilin 4
 hyaluronan-mediated motility receptor (RHAMM)
 gamma-aminobutyric acid (GABA) A receptor, alpha 1
 syntaxin 11
 crystallin, alpha B
 Consensus includes gb:AW839210 /FEA=EST /DB_XREF=gi:7930184 /DB_XREF=est:RC3-LT0023-200100-012-e09 /UG=Hs.278648 Homo sapiens cDNA FLJ14085 fis. t AW839210
 maternally expressed 3
 ribonuclease, RNase A family, k6
 prostaglandin D2 synthase (21kD, brain)
 Consensus includes gb:AL049285.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564M193 (from clone DKFZp564M193) /FEA=mRNA /DB_XREF=gi:4500045 /UG=Hs.302C AL049285
 defensin, beta 1
 superoxide dismutase 3, extracellular
 cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 6
 AF15q14 protein
 receptor tyrosine kinase-like orphan receptor 2
 aquaporin 1 (channel-forming integral protein, 28kD)
 Consensus includes gb:H15535 /FEA=EST /DB_XREF=gi:880355 /DB_XREF=est:ym27c01.s1 /CLONE=IMAGE:49385 /UG=Hs.52792 Homo sapiens mRNA: cDNA DKFZ: H15535
 homeo box C8
 type I transmembrane protein Fn14
 glutathione S-transferase M2 (muscle)
 butyrophilin, subfamily 3, member A3
 NADPH oxidase 1
 group III secreted phospholipase A2
 synaptotagmin 1
 tumor endothelial marker 8
 KIAA1107 protein
 cytochrome c oxidase subunit IV isoform 1
 spindlin-like
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12)
 KIAA1048 protein
 trans-phenyltransferase
 rat finger protein-like 1
 splicing factor, arginine/serine-rich 6
 translocation protein 1
 CD44 antigen (homing function and Indian blood group system)
 procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2
 chloride channel, calcium activated, family member 1
 D-aspartate oxidase
 paroxonase 1
 matrix metalloproteinase 27
 glycoprotein Ib (platelet), alpha polypeptide
 hypothetical protein FLJ20254
 like-glycosyltransferase
 Consensus includes gb:AL157490.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434F1011 (from clone DKFZp434F1011) /FEA=mRNA /DB_XREF=gi:7018535 /UG=Hs.27: AL157490
 cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6
 Consensus includes gb:AL359557.1 /DEF=Homo sapiens mRNA: cDNA DKFZp762O1415 (from clone DKFZp762O1415) /FEA=mRNA /DB_XREF=gi:8655612 /UG=Hs.30 AL359557
 fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism)
 KIAA0485
 cell division cycle 2-like 5 (cholinesterase-related cell division controller)
 Consensus includes gb:AU145003 /FEA=EST /DB_XREF=gi:11006524 /DB_XREF=est:AU145003 /CLONE=HEMBA1003598 /UG=Hs.287428 Homo sapiens cDNA FLJ111 AU145003
 gb:NM_020484.1 /DEF=Homo sapiens RAGE binding protein (AF011757), mRNA, /FEA=mRNA /GEN=AF011757 /PROD=RAGE binding protein /DB_XREF=gi:10048411 / NM_020484
 erythrocyte membrane protein band 4.9 (dema1n)
 phosphodiesterase 8A
 interleukin 1 receptor antagonist
 ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
 5-hydroxytryptamine (serotonin) receptor 1B
 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C
 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
 Consensus includes gb:AF043583.1 /DEF=Homo sapiens clone ASMmeg1-63 immunoglobulin lambda chain VJ region, (IGL) mRNA, partial cds. /FEA=mRNA /GEN=IGL /FEA=AF043583
 golgi associated, gamma adaptin ear containing, ARF binding protein 3
 Eeg-related gene member 2
 sdenyosucchnata lysse
 hypothetical protein FLJ10916
 NM_018271

hypothetical protein MGC5457	BC000988	1,28
otacornedin 1	NM_006334	1,28
solute carrier family 31 (copper transporters), member 2	NM_001860	1,28
WD repeat domain 12	NM_018256	1,28
LIM domain protein	BC003096	1,28
solute carrier family 17 (anion/sugar transporter), member 5	NM_012434	1,28
hypothetical protein from clone 24796	AF070598	1,28
disrupted in schizophrenia 1	NM_018662	1,28
putative DNA/chromatin binding motif	AA729218	1,28
FN5 protein	NM_020179	1,28
wingless-type MMTV integration site family, member 6	AI191771	1,28
DnaJ (Hsp40) homolog, subfamily B, member 5	AK023253	1,28
Mdm4, transformed 3T3 cell double minute 4, p53 binding protein (mouse)	NM_007393	1,28
cathepsin G	NM_001911	1,28
KIAA1117 protein	AB028040	1,28
botulin (collagen/tumorinogen domain-containing) 1	NM_002003	1,28
hypothetical protein MGC4883	NM_024514	1,28
nut-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	U19601	1,28
phospholipase C, beta 4	NM_000933	1,28
mitogen-activated protein kinase kinase 4	AA810268	1,28
NAD kinase	BE551347	1,28
scavenger box F1	NM_001451	1,28
ecdy LDL receptor; SREC-scavenger receptor expressed by endothelial cells	NM_003693	1,28
melanoma antigen, family C, 1	NM_005462	1,28
tryptophan 2,3-dioxygenase	NM_005651	1,28
hypothetical protein FLJ12988	NM_024999	1,28
inositol 1,4,5-triphosphate receptor, type 2	AB012610	1,28
solute carrier family 25 (mitochondrial carrier; peroxisomal membrane protein, 34kD), member 17	BC006557	1,28
heme oxygenase (decycling) 1	NM_002133	1,28
Cockayne syndrome 1 (classical)	NM_000082	1,28
deleted in azoospermia 3	NM_020364	1,28
homolog of rat orphan transporter v7-3	NM_018057	1,28
TNFRSF1A-associated via death domain	NM_003789	1,28
dermatopontin	AI146848	1,28
small muscle protein, X-linked	NM_014332	1,28
inhibitor of growth family, member 1	AW193656	1,28
olfactory receptor, family 1, subfamily D, member 2	U53583	1,28
DKFZP434A043 protein	NM_015396	1,28
cytochrome b-245, beta polypeptide (chronic granulomatous disease)	AI308863	1,28
angiotensin-like 4	NM_016109	1,28
hypothetical protein FLJ22202	NM_024883	1,28
PK506 binding protein 9 (63 kD)	AL050187	1,28
Consensus includes gb:AL049260.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564E233 (from clone DKFZp564E233). /FEA=mRNA /DB_XREF=gi:4500007 /UG=Hs.3020	AL049260	1,28
eukaryotic translation initiation factor 4 gamma, 3	NM_003760	1,28
dysferlin, axonemal, light intermediate polypeptide	NM_003462	1,28
SRY (sex determining region Y)-box 18	NM_018418	1,28
3-hydroxybutyrate dehydrogenase (heart, mitochondrial)	BC005844	1,28
serine/threonine kinase 38	AF034187	1,28
KIAA0328 protein	AI825212	1,28
uracil-DNA glycosylase	NM_003362	1,28
Consensus includes gb:U95737 /DEF=Human Chromosome 16 BAC clone CIT987SK-A-388D4 /FEA=mRNA /DB_XREF=gi:3417290 /UG=Hs.177961 Human Chromosome	U95737	1,28
differentiation-related protein dif13	NM_016380	1,28
transmembrane protease, serine 5 (spinesin)	NM_030770	1,28
growth factor receptor-bound protein 14	NM_004490	1,28
proline and glutamic acid rich nuclear protein	BC002875	1,28
hypothetical protein FLJ13081	NM_024634	1,28
isocitrate dehydrogenase 2 (NADP+), mitochondrial	AI151428	1,28
phosphoinositid 3-phosphate-binding protein-2	BC000568	1,28
tachykinin receptor 3	NM_001059	1,28
C-reactive protein, pentraxin-related	NM_000587	1,28
membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2)	NM_005374	1,28
transmembrane 4 superfamily member 1	AI189753	1,28
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 2	NM_005935	1,28

hypothetical protein FLJ14126
 Consensus includes gb:AK021576.1 /DEF=Homo sapiens cDNA FLJ11514 f1s, clone HEMBA1002229, /FEA=mRNA /DB_XREF=gi:10432783 /UG=Hs.306602 Homo sapiens
 polyomavirus binding protein-interacting protein 1
 ELAV (embryonic lethal, abnormal vision, Orophiila)-like 2 (Hu antigen B)
 Consensus includes gb:AA131324 /FEA=EST /DB_XREF=gi:1693011 /DB_XREF=est:237b04.s1 /CLONE=IMAGE:503503 /UG=Hs.28264 Homo sapiens mRNA; cDNA DK-AA131324
 ras homolog gene family, member E
 DNA segment on chromosome X and Y (unique) 155 expressed sequence
 CCAAT/enhancer binding protein (C/EBP) delta
 Consensus includes gb:AK025194.1 /DEF=Homo sapiens cDNA: FLJ21541 f1s, clone COL06166, /FEA=mRNA /DB_XREF=gi:10437657 /UG=Hs.306784 Homo sapiens cDNA
 DKFZP568A0522 protein
 hypothetical protein FLJ23074
 myosin, heavy polypeptide 11, smooth muscle
 PAN2 protein
 astrotactin 2
 dystroglycan 1 (dystrophin-associated glycoprotein 1)
 general transcription factor IIA, 1 (37KD and 19KD subunits)
 potassium voltage-gated channel, Shab-related subfamily, member 1
 glycoprotein M6B
 Consensus includes gb:AL049285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F053 (from clone DKFZp564F053), /FEA=mRNA /DB_XREF=gi:4500013 /UG=Hs.71981 AL049285
 matrix metalloproteinase 12 (macrophage elastase)
 TGFB inducible early growth response
 Consensus includes gb:AU148255 /FEA=EST /DB_XREF=gi:1009776 /DB_XREF=est:AU148255 /CLONE=MAMMA1003031 /UG=Hs.287627 Homo sapiens cDNA FLJ14
 hypothetical protein FLJ10312
 zinc finger protein 165 (LIM domain)
 beta-galactose-3-O-sulfotransferase, 4
 PRO233 protein
 tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor)
 paired immunoglobulin-like receptor alpha
 atrophin-1 interacting protein 1; actinin receptor interacting protein 1
 hypothetical protein
 Consensus includes gb:X63966.1 /DEF=H sapiens synthetic gene for platelet-derived growth factor-BB, /FEA=mRNA (PROD=platelet-derived growth factor-BB /DB_XREF=X63966
 gb:UA1742.1 /DEF=Human nucleophosmin-ratidic acid receptor alpha fusion protein NPM-RAR long form mRNA, complete cds, /FEA=mRNA (PROD=nucleophosmin-rel
 UA1742
 hypothetical protein FLJ20718
 KIAA1466 protein
 oxysterol-binding protein-related protein 1
 adaptor-related protein complex 1, gamma 2 subunit
 myotubularin related protein 3
 hypothetical protein MGC3020
 ubiquitin-conjugating enzyme E2L 6
 Consensus includes gb:AL137475.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F0723 (from clone DKFZp434F0723), /FEA=mRNA /DB_XREF=gi:6808086 /UG=Hs.27
 angiotensin-like factor
 ret finger protein-like 3 antisense
 Misshapen/NIK-related kinase
 transcription termination factor, RNA polymerase II
 sialic acid binding Ig-like lectin 7
 ankyrin 1, erythrocytic
 zinc finger protein 133 (clone pHZ-13)
 hypothetical protein LOC37187
 hypothetical protein FLJ11506
 interferon gamma receptor 1
 serologically defined colon cancer antigen 1
 transcription factor 7 (T-cell specific, HMG-box)
 hypothetical protein MGC4293
 hypothetical protein FLJ20896
 KIAA0404 protein
 ferritin, light polypeptide
 thrombomodulin
 microsomal glutathione S-transferase 3
 X11L-binding protein 51
 hypothetical protein FLJ22002
 cytoskeleton associated protein 2
 endothelin converting enzyme 2
 Consensus includes gb:AU145806 /FEA=EST /DB_XREF=gi:11007327 /DB_XREF=est:AU145806 /CLONE=HEMBA1005894 /UG=Hs.286664 Homo sapiens cDNA FLJ11; AU145806

hypothetical protein FLJ20428	NM_017817	1.28
E74-like factor 5 (ets domain transcription factor)	AF115403	1.28
matrix metalloproteinase 14 (membrane-inserted)	NM_004995	1.28
hypothetical protein	AV700415	1.28
protein tyrosine phosphatase, receptor type, R	NM_002849	1.28
hypothetical protein MGC10902	NM_030570	1.28
homo box A10	BF792917	1.28
KIAA0261 protein	D87450	1.28
tripartite motif-containing 3	NM_006458	1.28
ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	AI140752	1.28
Consensus includes gb:AW056600 /FEA=EST:5934239 /DB_XREF=est:wx23h08.x1 /CLONE=IMAGE:2544543 /UG=Hs.18593 Homo sapiens cDNA: FLJ21	AW056600	1.28
peroxisome biogenesis factor 10	AB013818	1.28
SA (rat hypertension-associated) homolog	NM_005622	1.28
MAD, mothers against decapentaplegic homolog 2 (Drosophila)	AW151617	1.28
KIAA0934 protein	NZ2859	1.28
Consensus includes gb:AL122122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L098 (from clone DKFZp434L098). /FEA=mRNA /DB_XREF=gi:8102949 /UG=Hs.2745	AL122122	1.28
calpain 9 (mCL-4)	NM_008615	1.28
asporin (LRR class 1)	NM_017680	1.28
defensin, alpha 6, Paneth cell-specific	NM_001926	1.28
PC326 protein	NM_018442	1.27
nucleotide binding protein 1 (MinD homolog, E. coli)	NM_002484	1.27
zinc finger protein 14 (K0X 6)	NM_021030	1.27
transporter 1, ATP-binding cassette, sub-family B (MDR/TP)	NM_000593	1.27
Consensus includes gb:AL050032.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566F1224 (from clone DKFZp566F1224). /FEA=mRNA /DB_XREF=gi:4884272 /UG=Hs.30	AL050032	1.27
hypothetical protein FLJ20047	NM_017639	1.27
protease inhibitor 15	NM_015886	1.27
ryenodine receptor 3	NM_001036	1.27
retinal short-chain dehydrogenase/reductase retSDR2	NM_018245	1.27
LIM domain only 4	SC003600	1.27
Consensus includes gb:A1813483 /FEA=EST /DB_XREF=est:ty37e12.x1 /CLONE=IMAGE:2281294 /UG=Hs.324504 Homo sapiens mRNA: cDNA	AI813483	1.27
protein tyrosine phosphatase, receptor type, J	AB00462	1.27
KIAA0356 gene product	AB002354	1.27
transcription factor 7-like 2 (T-cell specific, HMG-box)	AI375916	1.27
T-cell acute lymphocytic leukemia 1	NM_003189	1.27
Consensus includes gb:AF218021.1 /DEF=Homo sapiens clone PP552 unknown mRNA. /FEA=mRNA /PROD=unknown /DB_XREF=gi:10441971 /UG=Hs.302127 Homo	AF218021	1.27
Wilms tumor 1	NM_024426	1.27
RAB2, member RAS oncogene family	AI743756	1.27
Consensus includes gb:N71739 /FEA=EST /DB_XREF=gi:1228451 /DB_XREF=est:wy52h11.r1 /CLONE=IMAGE:255809 /UG=Hs.237354 Homo sapiens cDNA FLJ14308	NM_024676	1.27
hypothetical protein FLJ22938	BE567977	1.27
KRAB box containing C2H2 type zinc finger bA526D8.4	NM_005766	1.27
FERM, RhoGEF (ARHGGEF) and pleckstrin domain protein 1 (chondrocyte-derived)	NM_004234	1.27
zinc finger protein 93 homolog (mouse)	AL050032	1.27
Consensus includes gb:AL050032.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566F1224 (from clone DKFZp566F1224). /FEA=mRNA /DB_XREF=gi:4884272 /UG=Hs.30	M29335	1.27
major histocompatibility complex, class II, DO alpha	AI340239	1.27
G-2 and S-phase expressed 1	NM_014044	1.27
DKFZP564G0222 protein	ZZ5426	1.27
mitogen-activated protein kinase kinase 5	NM_001128	1.27
gb:BC001957.1 /DEF=Homo sapiens, Similar to KIAA0144 gene product, clone MGC-781, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0144 gene product /	BC001957	1.27
Consensus includes gb:AK024315.1 /DEF=Homo sapiens cDNA FLJ14253 flt, clone OVARC1001376. /FEA=mRNA /DB_XREF=gi:10436666 /UG=Hs.300694 Homo	AK024315	1.27
Consensus includes gb:AL390157.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D179 (from clone DKFZp434D179). /FEA=mRNA /DB_XREF=gi:3368847 /UG=Hs.3065	AL390157	1.27
MCF 2 cell line derived transforming sequence	X13230	1.27
Consensus includes gb:AK021614.1 /DEF=Homo sapiens cDNA FLJ11552 flt, clone HEMBA1003021. /FEA=mRNA /DB_XREF=gi:10432830 /UG=Hs.306606 Homo	AK021614	1.27
collagen, type V, alpha 1	N30339	1.27
adaptor-related protein complex 1, gamma 1 subunit	NM_001128	1.27
lankyrase, TRF-1-interacting ankym-related ADP-ribose polymerase	AF082559	1.27
KIAA0318 gene product	AW297087	1.27
vasodilator-stimulated phosphoprotein	NM_003370	1.27
protein kinase (cAMP-dependent, catalytic) inhibitor alpha	NM_006823	1.27
hypothetical protein	BC005004	1.27
thymidine kinase 2, mitochondrial	BE895437	1.27
butyrophilin-like 3	AK025287	1.27
matrix metalloproteinase 8 (neutrophil collagenase)	NM_002424	1.27
BAI1-associated protein 2	NM_017451	1.27

transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
endothelin 3
KIAA0826 protein
TYRO3 protein tyrosine kinase
Consensus includes gb:AL117659.1 /DEF=Homo sapiens mRNA; cDNA DKFZp58610624 (from clone DKFZp58610624) /FEA=mRNA /DB_XREF=gi:5912252 /UG=Hs.3086 /AL117659
calcium channel, voltage-dependent, alpha 11 subunit
cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)
tumor necrosis factor (ligand) superfamily, member 4 (tax-transcriptionally activated glycoprotein 1, 34kD)
Consensus includes gb:AK000995.1 /DEF=Homo sapiens cDNA FLJ10133.1fa, clone HEMBA1003087 /FEA=mRNA /DB_XREF=gi:7021998 /UG=Hs.272209 Homo sapiens
Consensus includes gb:AI191118 /FEA=EST /DB_XREF=gi:3742327 /DB_XREF=est:q16c05.x1 /CLONE=IMAGE:1739336 /UG=Hs.222015 ESTs
stromatin (EBP72)-like 1
Consensus includes gb:AL031686 /DEF=Human DNA sequence from clone 981L23 on chromosome 20q12.1-13.2. Contains a Krueppel type zinc-finger protein pseudogen
ribonuclease, RNase A family, 4
cardiotrophin-like cytokine: neurotrophin-1/B-cell stimulating factor-3
hypothetical protein FLJ10598
leukocyte membrane antigen
LIM domain binding 3
KIAA0061 protein
Consensus includes gb:R25849 /FEA=EST /DB_XREF=gi:781984 /DB_XREF=est:Y42a11.r1 /CLONE=IMAGE:132380 /UG=Hs.217305 ESTs
methylmalonyl Coenzyme A mutase
TBP-like 1
KIAA0671 gene product
KIAA0878 protein
Consensus includes gb:AF238870.1 /DEF=Homo sapiens clone GLSH-3 similar to gliadin mRNA sequence. /FEA=mRNA /DB_XREF=gi:8272577 /UG=Hs.275706 Homo sapiens
myelin-associated oligodendrocyte basic protein
FGFR1 oncogene partner
protocadherin gamma subfamily A, 1
Consensus includes gb:AX023116.1 /DEF=Homo sapiens cDNA FLJ13054.1fa, clone NT2RP3001527, highly similar to Human Sp140 protein (Sp140) mRNA. /FEA=mRNA /DB_XREF=gi:8272577 /UG=Hs.275706 Homo sapiens
hemochromatosis
light junction protein 1 (zone occludens 1)
connexin-36
Consensus includes gb:AL353942.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L23121 (from clone DKFZp761L23121) /FEA=mRNA /DB_XREF=gi:7669981 /UG=Hs.275706 Homo sapiens
lipocalin 2 (oncoferrin 24p3)
hypothetical protein FLJ11467
Consensus includes gb:AF107846 /DEF=Homo sapiens neuroendocrine-specific Golgi protein p55 (XLalpha) gene, exon XL2 and complete cds /FEA=mRNA /DB_XREF=gi:7669981 /UG=Hs.275706 Homo sapiens
RAB interacting factor
ralA binding protein 1
plexin A2
interleukin-1 homolog 1
ectonucleoside triphosphate diphosphohydrolase 2
estrogen-related receptor gamma
Consensus includes gb:AI375002 /FEA=EST /DB_XREF=gi:4174992 /DB_XREF=est:ts4b11.x1 /CLONE=IMAGE:2047869 /UG=Hs.272014 ESTs. Weakly similar to ALU3 /AI375002
collagen, type VI, alpha 1
lymphocyte antigen 95 (activating NK-receptor, NK-p44)
KIAA0984 protein
synaptin 6
TAF7 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 55 kD
KIAA0276 protein
serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1
X03453 Bacteriophage P1 cre recombinase protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)
DKFZP564O123 protein
opiate receptor-like 1
chloride channel, calcium activated, family member 2
v-jes-1 Y-enaguchi sarcoma viral oncogene homolog 1
coagulation factor IX (plasma thromboplastin component, Christmas disease, hemophilia B)
integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor)
nuclear factor I/C (CCAAT-binding transcription factor)
lipocalin-interacting membrane receptor
hypothetical protein FLJ10709
Consensus includes gb:AU152579 /FEA=EST /DB_XREF=gi:11014100 /DB_XREF=est:AU152579 /CLONE=NT2RP3001232 /UG=Hs.208931 Homo sapiens cDNA FLJ130 /AU152579
Consensus includes gb:R73554 /FEA=EST /DB_XREF=gi:8417586 /DB_XREF=est:Y94e05.x1 /CLONE=IMAGE:156418 /UG=Hs.103391 Human insulin-like growth factor bi R73554
phosphodiesterase A2, group IIA (platelets, synovial fluid)
HCF-binding transcription factor Zhangfei

DKFZP434F1735 protein	NM_015590	1,27
KIAA1109 protein	AL137254	1,27
keratinophilin alpha 2 (RAG cohort 1, importin alpha 1)	NM_002286	1,27
Consensus includes gb:AU147182 /FEA=EST /DB_XREF=gi:11008703 /DB_XREF=au:AU147182 /CLONE=HEWBB1002623 /UG=Hs.301197 Homo sapiens cDNA FLJ121	NM_002286	1,27
Consensus includes gb:AU147182 /FEA=EST /DB_XREF=gi:11008703 /DB_XREF=au:AU147182 /CLONE=HEWBB1002623 /UG=Hs.301197 Homo sapiens cDNA FLJ121	AF022231	1,27
conserved gene amplified in osteosarcoma	AY004175	1,27
phosphoinositide-specific phospholipase C-beta 1	NM_024763	1,27
hypothetical protein FLJ23129	NM_017715	1,27
hypothetical protein FLJ20216	NM_049259	1,27
Consensus includes gb:AL049259.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E193 (from clone DKFZp564E193) /FEA=mRNA /DB_XREF=gi:4500005 /UG=Hs.11051	AL072238	1,27
adenylosuccinate lyase	NM_014097	1,27
PRO1693 protein	NM_004681	1,27
eukaryotic translation initiation factor 1A, Y chromosome	BE966236	1,27
ribonucleotide reductase M2 polypeptide	BC000419	1,27
catechol-O-methyltransferase	A457120	1,27
phosphoribosyl pyrophosphate amidotransferase	AF020543	1,27
palmitoyl-protein thioesterase 2	NM_000672	1,27
alcohol dehydrogenase 6 (class V)	AA837026	1,27
Consensus includes gb:AA837026 /FEA=EST /DB_XREF=gi:2912225 /DB_XREF=est:cg09c12.s1 /CLONE=IMAGE:1439350 /UG=Hs.163549 ESTs	ST0123	1,27
low density lipoprotein receptor (familial hypercholesterolemia)	NM_000123	1,27
excision repair cross-complementing rodent repair deficiency, complementation group 5 (xeroderma pigmentosum, complementation group G (Cockayne syndrome))	AF053356	1,27
erythropoietin	R94844	1,27
Consensus includes gb:R94844 /FEA=EST /DB_XREF=gi:970039 /DB_XREF=est:yt42a12.1 /CLONE=IMAGE:199430 /UG=Hs.306542 Homo sapiens version Vini isoform	J04798	1,27
prothymosin, alpha (gene sequence 28)	NM_016308	1,27
hypothetical protein FLJ11042	NM_002278	1,27
keratin, hair acidic, 2	BC005253	1,27
Kelch motif containing protein	NM_022453	1,27
ring finger protein 25	NM_018314	1,27
hypothetical protein FLJ11088	NM_002637	1,27
phosphorylase kinase, alpha 1 (muscle)	U84138	1,27
RAD51-like 1 (S. cerevisiae)	BF590317	1,27
HYA22 protein	NM_020664	1,27
2,4-dienoyl CoA reductase 2, peroxisomal	AL523320	1,27
hypothetical protein FLJ10097	NM_018138	1,27
hypothetical protein FLJ10560	M96843	1,27
striated muscle contraction regulatory protein	A1728958	1,27
titin 2	AJ275355	1,27
Consensus includes gb:AJ275355 /DEF=Homo sapiens partial IGTVH3 V3-20 gene for immunoglobulin heavy chain V region, case 1, clone 2 /FEA=CDS /DB_XREF=gi:757; A	NM_004491	1,27
glucocorticoid receptor DNA binding factor 1	NM_002550	1,27
offactory receptor, family 3, subfamily A, member 1	NM_007056	1,27
calpain 11	NM_002054	1,27
glucagon	NM_008225	1,27
phospholipase C, delta 1	A1743090	1,27
hypothetical protein FLJ23138	BG435463	1,27
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	NM_018891	1,27
laminin, gamma 2 (niclin (100KD), laminin (105KD), BM600 (100KD), Herlitz junctional epidermolysis bullosa))	BC006403	1,27
NCK adaptor protein 1	NM_003893	1,27
LIM domain binding 1	NM_015148	1,27
PAS-serine/threonine kinase	NM_018725	1,27
IL-17B receptor	D88435	1,27
cyclin G associated kinase	NM_007202	1,27
A kinase (PRKA) anchor protein 10	AL049223	1,27
secretory carrier membrane protein 1	NM_004311	1,27
ADP-ribosylation factor-like 3	Y15816	1,27
collagen, type I, alpha 1	NM_005822	1,27
Down syndrome critical region gene 1-like 1	BC530461	1,27
KIAA0582 protein	NM_004593	1,27
RAB5C, member RAS oncogene family	NM_014852	1,27
KIAA0945 protein	AW304174	1,27
chitinase, di-N-acetyl-	NM_030613	1,27
hypothetical protein FLJ21628	BC002977	1,27
mel transforming oncogene (derived from cell line NK14)- RAB8 homolog	AL136727	1,27
RAB8C, member RAS oncogene family	AB501145	1,27
phosphodiesterase 4D interacting protein (myomegalin)		

melanoma antigen, family A, 4
cyclin-E binding protein 1
activin A receptor, type I
winged-helix nude
geminin
hypothetical protein FLJ10595
Consensus includes gb:AL0409593 /DEF=Human DNA sequence from clone RP4-654A7 on chromosome 20. Contains the gene for phospholipase C-beta-1 (KIAA0581, 1-p AL049593)
Consensus includes gb:AL137798 /DEF=Human DNA sequence from clone RPS-1182A14 on chromosome 1 Contains part of a gene similar to rat Espin, a pseudogene
polydendylate binding protein-interacting protein 1
MAD, mothers against decapentaplegic homolog 9 (Drosophila)
KIAA0172 protein
BSP-like protein 2
v-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog
aldehyde dehydrogenase 3 family, member A2
growth factor receptor-bound protein 10
hypothetical protein DKFZP434K1210
transmembrane protein vezatin
zinc finger protein 143 (clone pHz-1)
DKFZP586F1018 protein
hypothetical protein FLJ11149
proprotein convertase subtilisin/kexin type 1
transducer of ERBB2, 2
phospholipase C, epsilon
KIAA0513 gene product
Consensus includes gb:AK000185.1 /DEF=Homo sapiens cDNA FLJ20178 fts, clone COL09990. /FEA=mRNA /DB_XREF=gi:7020104 /UG=Hs.306389 Homo sapiens cDN
Huntingtin interacting protein H
DKFZP434M154 protein
deleted in lymphocytic leukemia, 2
Consensus includes gb:BC002465.1 /DEF=Homo sapiens, clone IMAGE:3344506. mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3344506) /DB_XI
Mad4 homolog
TU3A protein
Consensus includes gb:AF688610 /FEA=EST /DB_XREF=gi:4827918 /DB_XREF=est:y56a05.x5 /CLONE=IMAGE:162224 /UG=Hs.32043 ESTs
surfactant protein B
growth factor receptor-bound protein 10
KIAA0924 protein
dynein, cytoplasmic, intermediate polypeptide 1
Wolf-Hirschhorn syndrome candidate 1-like 1
Consensus includes gb:AF055018.1 /DEF=Homo sapiens clone 24442 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005745 /UG=Hs.139137 Homo sapiens clone 24442
PXR2b protein
Consensus includes gb:AL050204.1 /DEF=Homo sapiens mRNA; cDNA DKFZ586F1223 (from clone DKFZ586F1223). /FEA=mRNA /DB_XREF=gi:4884443 /UG=Hs.28
KIAA1005 protein
lethal (3) malignant brain tumor (3)mbt protein (Drosophila) homolog
protein (pepity-proty clatrants isomerase) NIMA-interacting, 4 (parvulin)
RAD23 homolog B (S. cerevisiae)
hypothetical protein FLJ20456
solute carrier family 13 (sodium/sulfate symporters), member 4
zinc finger protein 83 (HPF1)
BLUB1 budding uninhibited by benzimidazoles 1 homolog (yeast)
Consensus includes gb:AL105684.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 27080. /FEA=mRNA /DB_XREF=gi:5689805 /UG=Hs.306329 I AL109684
Consensus includes gb:X55989 /DEF=Human ECRP gene for eosinophil cationic related protein /FEA=mRNA /DB_XREF=gi:22940 /UG=Hs.166037 Human ECRP gene for X55989
Kruppel-like factor 1 (erythroid)
T-cell leukemia/lymphoma 6
weakly similar to glutathione peroxidase 2
insulin-like growth factor 1 (somatomedin C)
FK506 binding protein 8 (38kD)
flavin containing monooxygenase 3
hypothetical protein MGCA701
atriatin, calmodulin binding protein
pyruvate dehydrogenase kinase, isoenzyme 2
KIAA0599 protein
growth arrest-specific 7
hypothetical protein FLJ12975
Consensus includes gb:AL109681.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 112333. /FEA=mRNA /DB_XREF=gi:5689785 /UG=Hs.226017 AL109681

228

oxygen regulated protein (150kD)	NM_006389	1.26
transcription factor (p38 interacting protein)	A4889653	1.26
protein with polyglutamine repeat; calcium (ca2+) homeostasis endoplasmic reticulum protein	A3737979	1.26
KIAA1919 protein	AK000168	1.26
melanoma antigen, family A, 8	BC002455	1.26
cyclodrome b-5	M22978	1.26
CGI-75 protein	NM_016020	1.26
prostaglandin E receptor 2 (subtype EP2), 53kD	NM_000956	1.26
Consensus includes gb:AA195336 /FEA=EST /DB_XREF=gi:1791527 /DB_XREF=est:zp95d04.s1 /CLONE=IMAGE:628327 /UG=Hs.82719 Homo sapiens mRNA; cDNA DI AA195936	NM_000956	1.26
KIAA1324 protein	AB037745	1.26
ubiquitin specific protease 12	A820101	1.26
hypothetical protein FLJ10450	NM_018095	1.26
TAF10 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 30 kD	NM_006284	1.26
hypothetical protein FLJ12328	NM_024787	1.26
Consensus includes gb:AV700338 /FEA=EST /DB_XREF=gi:10302309 /DB_XREF=est:AV700338 /CLONE=GKCAMH12 /UG=Hs.286243 ESTs	AV700338	1.26
Kelch-like ECH-associated protein 1	NM_012289	1.26
acetylserotonin O-methyltransferase	NM_004043	1.26
lipase, hepatic	NM_000236	1.26
homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease	BC000747	1.26
TNF receptor-associated factor 1	NM_005658	1.26
renal tumor antigen	NM_014226	1.26
hypothetical protein FLJ22318	NM_022762	1.26
potassium voltage-gated channel, KQT-like subfamily, member 1	AF051426	1.26
lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD)	NM_005565	1.26
sialic acid binding Ig-like lectin 6	NM_001245	1.26
Consensus includes gb:AV468201 /FEA=EST /DB_XREF=gi:7038307 /DB_XREF=est:he34a04.x1 /CLONE=IMAGE:2920878 /UG=Hs.91375 Human clone 23614 mRNA	NM_001245	1.26
nuclear phosphoprotein similar to S. cerevisiae PWP1	BE796924	1.26
RNA isopentenylpyrophosphate transferase	AF116679	1.26
camitine palmitoyltransferase L, liver	NM_001876	1.26
Consensus includes gb:AK024871.1 /DEF=Homo sapiens cDNA: FLJ21218 fs, clone COL00537, /FEA=mRNA /DB_XREF=gi:10437282 /UG=Hs.306712 Homo sapiens cDNA AK024871	NM_001876	1.26
trophinin	NM_016157	1.26
ADP-ribosyltransferase 3	U47054	1.26
reticulocalbin 1, EF-hand calcium binding domain	NM_002901	1.26
EH-domain containing 3	NM_014600	1.26
Consensus includes gb:AIB84814 /FEA=EST /DB_XREF=gi:5569978 /DB_XREF=est:w85e05.x1 /CLONE=IMAGE:2431664 /UG=Hs.21964 ESTs	NM_014600	1.26
melanoma antigen, family A, 12	AIB84814	1.26
gb.L14457.1 /DEF=Human Ig rearranged kappa-chain gene V-J-region, complete cds, /FEA=mRNA /DB_XREF=gi:398491 /FL=gb.L14457.1	U82871	1.26
amine oxidase, copper containing 2 (retine-specific)	L14457	1.26
histamine N-methyltransferase	NM_009590	1.26
hypothetical protein PRO1777	NM_006895	1.26
KIAA0671 gene product	NM_018805	1.26
Consensus includes gb:AU145203 /FEA=EST /DB_XREF=gi:11006724 /DB_XREF=est:AU145203 /CLONE=HEMBA1004193 /UG=Hs.300897 Homo sapiens cDNA FLJ111 AU145203	NM_018805	1.26
tenascin XB	AL136896	1.26
AU RNA binding protein/enzyme-Coenzyme A hydratase	BE044614	1.26
secreted phosphoprotein 2, 24kD	NM_001698	1.26
fibroblast growth factor 23	NM_006944	1.26
zinc finger protein 313	NM_020638	1.26
hypothetical protein MGC5528	AL031685	1.26
annexin A7	L11372	1.26
megakaryocyte-enhanced gene transcript 1 protein	NM_004034	1.26
olfactory receptor, family 32, subfamily A, member 1	NM_021246	1.26
KIAA0196 gene product	NM_012375	1.26
phosphoglycerate kinase 1	NM_014846	1.26
Consensus includes gb:AF090101.1 /DEF=Homo sapiens clone IMAGE 30714, /FEA=mRNA /DB_XREF=gi:4063636 /UG=Hs.177837 Homo sapiens clone IMAGE 30714	NM_014846	1.26
G protein-coupled receptor 39	AL121974	1.26
kalikrein B, plasma (Fletcher factor) 1	AF090101	1.26
Consensus includes gb:AU147194 /FEA=EST /DB_XREF=gi:11008715 /DB_XREF=est:AU147194 /CLONE=HEMBA1002684 /UG=Hs.300978 Homo sapiens cDNA FLJ12 AU147194	AF090101	1.26
ADP-ribosylation factor 3	NM_000892	1.26
calcium/calmodulin-dependent protein kinase (CaM kinase) II alpha	BQ341806	1.26
zinc finger protein 177	AW131813	1.26
Fancx1 anemia, complementation group G	NM_003451	1.26
tumor necrosis factor receptor superfamily, member 4	AC004472	1.26
olfactory receptor, family 12, subfamily D member 3	NM_003327	1.26
	NM_030959	1.26

zinc finger protein 286
hypothetical protein similar to preferentially expressed antigen of melanoma
erythropoietin receptor
SMC4 structural maintenance of chromosomes 4-like 1 (yeast)
membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific)
putative chemokine receptor; GTP-binding protein
KIAA0408 gene product
KIAA1750 protein
sperm acrosome associated 1
egrin
desmoolin 2
docking protein 1, 62KD (downstream of tyrosine kinase 1)
hypothetical protein MGC2827
matrix metalloproteinase 1 (interstitial collagenase)
dual specificity phosphatase 10
solute carrier family 2 (facilitated glucose transporter), member 10
hypothetical protein FLJ11017
KIAA0535 gene product
immunoglobulin mu binding protein 2
immunoglobulin lambda locus
UDP glycosyltransferase 2 family, polypeptide B15
kinesin family member 1C
Consensus includes gb:BF939689 /FEA=EST /DB_XREF=gi:2357100 /DB_XREF=est:nac80b04.x1 /CLONE=IMAGE:34-0575 /UG=Hs.159629 myosin IXB
DKF ZP564M1462 protein
KIAA1079 protein
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 12
chloride channel 2
Consensus includes gb:AK025072.1 /DEF=Homo sapiens cDNA: FLJ21419 fls, clone COL04084 /FEA=mRNA /DB_XREF=gi:10437512 /UG=Hs.306759 Homo sapiens cDNA: AK025072
radial spokehead-like 1
matrix metalloproteinase 23A
pellino homolog 2 (Drosophila)
brain abundant, membrane attached signal protein 1
laminin, gamma 1 (formerly LAMB2)
contactin 1
met proto-oncogene (hepatocyte growth factor receptor)
hypothetical protein FLJ10895
myelin proteolipid protein-like protein
hypothetical protein MGC13523
phorbolin (similar to spolliprotein B mRNA editing protein)
nect (homologous to the EG-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1
deoxyribonuclease II, lysosomal
mitochondrial elongation factor G
translin
syndecan binding protein (synterin)
TEK tyrosine kinase, endothelial (venous malformations, multiple cutaneous and mucosal)
chromobox homolog 8 (Pc class homolog, Drosophila)
Chediak-Higashi syndrome 1
solute carrier family 12 (potassium/chloride transporters), member 4
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3
dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)
collagen, type IX, alpha 1
I-complex 10 (a murine Icp homolog)
calsequestrin 2 (cardiac muscle)
HERV-H LTR-associating 1
transducin-like enhancer of split 1 (E(spl) homolog, Drosophila)
Consensus includes gb:AF070587.1 /DEF=Homo sapiens clone 24741 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387960 /UG=Hs.25770 Homo sapiens clone 24741, AF070587
activity-regulated cytoskeleton-associated protein
periplakin
suppressor of variegation 3-9 (Drosophila) homolog 2; hypothetical protein FLJ23414
hBOIT for potent brain type organic ion transporter
uncoupling protein 1 (mitochondrial, proton carrier)
Consensus includes gb:AU150943 /FEA=EST /DB_XREF=gi:11012464 /DB_XREF=est:AU150943 /CLONE=NT2R2P2003984 /UG=Hs.66762 Homo sapiens mRNA: cDNA (AU150943)
minor histocompatibility antigen HA-1

S-phase response (cyclin-related)
keratin, hair, acidic, 4
apoptosis signaling protein, nonapoptotic homolog (mouse)
Bacillus subtilis REF=M24537 /DEF=AB subtilis p8b, pheA genes corresponding to nucleotides 2897-3200 of M24537 /LEN=1409 (-5, -M, -3 represent transcript regions 5
M24537
potassium inwardly-rectifying channel, subfamily J, member 12
fetuin B
carboxylesterase 2 (intestine, liver)
arginase, liver
DC11 protein
KIAA0265 protein
Consensus includes gb:F09448 /FEA=EST /DB_XREF=gi:681981 /DB_XREF=est:HSC31G032 /CLONE=c-31g03 /UG=Hs.12840 Homo sapiens germline mRNA sequence
F09448
ata variant gene 3
apical protein-like (Xenopus laevis)
caspase 6, apoptosis-related cysteine protease
transmembrane 9 superfamily member 1
B-cell CLL/lymphoma 2
c-myc proto-oncogene tyrosine kinase
KIAA0557 protein
Consensus includes gb:AW452357 /FEA=EST /DB_XREF=gi:6993133 /DB_XREF=est:U1-H-B13-ai1-d-03-Q-U1.s1 /CLONE=IMAGE:3068213 /UG=Hs.27373 Homo sapiens
M12874
estrogen receptor 1
Consensus includes gb:BE217875 /FEA=EST /DB_XREF=gi:8905193 /DB_XREF=est:hv31a03.x1 /CLONE=IMAGE:3174988 /UG=Hs.12311 Homo sapiens clone 23570 m
BE217875
sodium channel, voltage-gated, type II, beta polypeptide
pyruvate carboxylase
secreted ficolin-related protein 4
Consensus includes gb:AK000645.1 /DEF=Homo sapiens cDNA FLJ20638 fls. clone KAT02982, highly similar to SUI5_RABIT SUCRASE-ISOMALTASE /FEA=mRNA ID
AK000645
zinc finger protein 288
EGF-containing fibulin-like extracellular matrix protein 2
transglutaminase 3 (E polypeptide, protein-glutamine-gamma-glutamyltransferase)
exostoses (multiple) 1
intracellular hyaluronan-binding protein
hypothetical protein MGC3046
POU transcription factor
solute carrier family 8 (sodium-calcium exchanger), member 2
solute carrier family 22 (organic anion transporter), member 7
cytidine deaminase
microtubule-associated protein 7
Consensus includes gb:AB051440.1 /DEF=Homo sapiens mRNA for KIAA1653 protein, partial cds. /FEA=mRNA /GEN=KIAA1653 /PROD=KIAA1653 protein /DB_XREF=gi
AB051440
mitogen-activating protein kinase kinase kinase 2
progestin induced protein
gb:NM_017910.2 /DEF=Homo sapiens hypothetical protein FLJ20628 (FLJ20628), mRNA. /FEA=mRNA /GEN=FLJ20628 /PROD=hypothetical protein FLJ20628 /DB_XREF=gi
NM_017910
factor H-related protein 5
splicing factor, arginine/serine-rich 7 (35kD)
hypothetical protein FLJ22041 similar to FK506 binding proteins
cholinergic receptor, nicotinic, alpha polypeptide 9
Consensus includes gb:AA898858 /FEA=EST /DB_XREF=gi:3015728 /DB_XREF=est:ak23d05.s1 /CLONE=IMAGE:1406793 /UG=Hs.118567 ESTs
potassium voltage-gated channel, shaker-related subfamily, beta member 1
Rhesus blood group, CcEa antigens
small nuclear RNA activating complex, polypeptide 1, 43kD
mucin 16
cyclin-dependent kinase inhibitor 1C (p57, Kip2)
DKFZP434F122 protein
Consensus includes gb:AF339785.1 /DEF=Homo sapiens clone IMAGE:1963178, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507323 /UG=Hs.270266 Homo sapiens
cytochrome P450, subfamily 48 (cholesterol 24-hydroxylase)
hypothetical protein FLJ12810
hypothetical protein FLJ13511
protein tyrosine phosphatase, receptor type, N polypeptide 2
hormone box A5
soluble adenylyl cyclase
T-box and WD-40 domain protein 3
cytochrome b-561
Htn-1
Consensus includes gb:BF739971 /FEA=EST /DB_XREF=gi:12066647 /DB_XREF=est:7a41107.x1 /CLONE=IMAGE:3576732 /UG=Hs.98416 ESTs
BF739971

discoidin domain receptor family, member 1
 period homolog 2 (Drosophila)
 membrane-spanning 4-domains, subfamily A, member 5
 DKEZP434C153 protein
 zinc finger protein, X-linked
 HEAT-like repeat-containing protein
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1
 U8 snRNA-associated Sm-like protein
 5-methyltetrahydrofolate-homocysteine methyltransferase
 Consensus includes gb:AU147799 /FEA=EST /DB_XREF=gi:11009320 /DB_XREF=est:AU147799 /CLONE=MAMMA1001744 /UG=Hs.288467 Homo sapiens cDNA FLJ12 AU147799
 SMT3 suppressor of mid two 3 homolog 2 (yeast)
 histamine H4 receptor
 hypothetical protein FLJ14050
 prostate stem cell antigen
 adaptor-related protein complex 1, sigma 2 subunit
 uradipropionase, beta
 keratin, hair, acidic 7
 hypothetical protein FLJ21439
 neuropilin
 hypothetical protein FLJ22558
 DKFZP586M0822 protein
 chondroitin sulfate proteoglycan 2 (versican)
 KIAA1071 protein
 KIAA1232 protein
 Consensus includes gb:AK024945.1 /DEF=Homo sapiens cDNA: FLJ21292 fs, clone COL01969, /FEA=mRNA /DB_XREF=gi:10437367 /UG=Hs.306732 Homo sapiens cd
 v-mpl myeloblastosis viral oncogene homolog (avian)
 origin recognition complex, subunit 6 (yeast homolog)-like
 seizure related 8 homolog (mouse)-like
 CGI-142
 ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) 42kD
 hypothetical protein
 IL2-inducible T-cell kinase
 Indian hedgehog homolog (Drosophila)
 estrogen-related receptor gamma
 Cluster ind. AL109698:Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 26539 /cds=UNKNOWN /gb=AL109698 /gi=5689808 /ug=Hs.8065 /len=2035
 lactate dehydrogenase A, like
 regulator of G-protein signalling 11
 histone deacetylase 8
 immunoglobulin kappa constant
 downstream neighbor of SON
 KIAA1128 protein
 optic atrophy 1 (autosomal dominant)
 KIAA0703 gene product
 hypothetical protein
 hypothetical protein FLJ21135
 TM7SF for transmembrane molecule with thrombospondin module
 KH-type splicing regulatory protein (FUSE binding protein 2)
 caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
 metallothionein 1E (functional)
 receptor (calcitonin) activity modifying protein 3
 myofibrin coreceptor synthesis 3
 ocular albinism 1 (Nethership-Falls)
 KIAA0663 protein
 KIAA0744 protein
 thymidine repeat containing 3
 dysprosophilin, beta
 phosphodiesterase 8H, cGMP-specific, cone, gamma
 phosphoinositide 3-phosphate-binding protein-2
 gb:BC006408.1 /DEF=Homo sapiens, Similar to zinc finger protein 254, clone MGC:10544, mRNA, complete cds. /FEA=mRNA /PROD=Similar to zinc finger protein 254 /ID BC006408
 soluble carrier family 23 (nucleoside transporters), member 1

mitochondrial capsule selenoprotein 1.25 NM_030663

chromosome 11 open reading frame 11 1.25 AB014559

protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform 1.25 NM_004156

protocadherin gamma subfamily C, 3 1.25 AF152524

RIG-like 5-6 1.25 NM_015682

gb:BC006441.1 /DEF=Homo sapiens. Similar to RNA polymerase I transcription factor RRN3, clone MGC:13169, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RN 1.25 BC006441

poly(ADP-ribose) polymerase 2 1.25 AW103422

hypothetical protein DKFZp762E1312 1.25 NM_018410

distal-less homeobox 4 1.25 NM_001934

EphB2 1.25 AF025304

Consensus includes gb:AV741657 /FEA=EST /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1 1.25 AV741657

chromodomain helicase DNA binding protein 4 1.25 NM_001273

prostaglandin D2 receptor (DP) 1.25 AI460323

cell division cycle 25C 1.25 AF277724

TAK1-binding protein 2 1.25 AL117407

hypothetical protein FLJ20366 1.25 NM_017786

mitogen-activated protein kinase-activated protein kinase 5 1.25 NM_003668

gb:AF116659.1 /DEF=Homo sapiens PRO1412 mRNA, complete cds. /FEA=mRNA /PROD=PRO1412 /DB_XREF=gi:7959816 /UG=Hs.321151 Homo sapiens PRO1412 m 1.25 AF116659

heterogeneous nuclear ribonucleoprotein L 1.25 X16135

KIAA0427 gene product 1.25 NM_014772

keratin 4 1.25 X07895

leucine-zipper-like transcriptional regulator, 1 1.25 NM_008767

insulin promoter factor 1, homeodomain transcription factor 1.25 U35632

solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7 1.25 NM_020309

NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) 1.25 NM_004549

p53-induced protein 1.25 NM_006034

hypothetical protein 1.25 NM_013341

Consensus includes gb:AW242315 /FEA=EST /DB_XREF=gi:6576069 /DB_XREF=IMAGE:2692055 /UG=Hs.27860 Homo sapiens mRNA; cDNA 1.25 AW242315

peanut-like 2 (Orsophilis) 1.25 AF176379

interleukin 18 binding protein 1.25 NM_005699

hypothetical protein FLJ10312 1.25 NM_030672

hypothetical protein 1.25 NM_016629

KIAA0171 gene product 1.25 NM_014866

Wilms tumor associated protein 1.25 NM_015855

microfilament-associated protein 2 1.25 AL049569

hypothetical protein MGC3047 1.25 AW888223

Consensus includes gb:AL049690 /DEF=Human DNA sequence from clone 1080K8 on chromosome 20p12.1-13 Contains a pseudogene similar to 40S ribosomal protein 1.25 AL049690

hypothetical protein 1.25 W72694

amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2) 1.25 AI141541

Consensus includes gb:AL137495.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C2331 (from clone DKFZp434C2331). /FEA=mRNA /DB_XREF=gi:5808119 /UG=Hs.30 1.25 AL137495

SRY (sex determining region Y)-box 11 1.25 AI360875

KIAA0578 protein 1.25 AL117532

chromosome 1 open reading frame 1 1.25 NM_001213

cholesterol receptor, muscarinic 2 1.25 NM_000739

ATPase, H+ transporting, lysosomal (vacuolar proton pump), alpha polypeptide, 70kD, isoform 1 1.25 AF113129

hypothetical protein FLJ10849 1.25 AI539361

protein tyrosine phosphatase, non-receptor type 22 (lymphoid) 1.25 NM_012411

reticulon 4 1.25 AF333338

fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome) 1.25 NM_023110

centaurin, beta 1 1.25 NM_014716

myeloproliferative leukemia virus oncogene 1.25 M90103

eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD) 1.25 NM_003750

adenomedullin 1.25 NM_001124

olfactory receptor, family 7, subfamily E, member 12 pseudogene 1.25 AF065854

major histocompatibility complex, class II, DQ beta 1 1.25 AW276186

Fe fragment of IgG low affinity Ila receptor for (CD32) 1.25 NM_021642

pregnancy specific beta-1-glycoprotein 3 1.25 NM_021016

protein C (inactivator of coagulation factors Va and VIIIa) 1.25 NM_000312

solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 1.25 AF064484

DEAD-box protein 1.25 NM_018665

adaptor-related protein complex 1, gamma 1 subunit 1.25 BF752277

KIAA0903 protein 1.25 BF116032

PDZ domain containing guanine nucleotide exchange factor(GEF)1 1.25 NM_014247

protein tyrosine phosphatase, non-receptor type 2 1.25
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease) 1.25
bradykinin receptor B1 1.25
G protein-coupled receptor 2 1.25
PBX/invited 1 homeobox 2 1.25
Cluster Ind. N53536;yz26h0.1.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:284209 /clone_end=3 /gb=N53536 /gi=1194702 /ug=Hs.33540 /len=543 1.25
ring finger protein 1 1.25
adenosine monophosphate deaminase (isoform E) 1.25
hypothetical protein FLJ22127 1.25
ubiquitin associated protein 1.25
lymphoid blast crisis oncogene 1.25
hypothetical protein FLJ22955 1.25
Mid-1-related chloride channel 1 1.25
glycerol kinase pseudogene 1 1.25
gb:BC000077.1 /DEF=Homo sapiens, Similar to KIAA0377 gene product, clone MGC:3268, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0377 gene product 1.25
a disintegrin and metalloproteinase domain 21 1.25
hypothetical protein FLJ23093 1.25
mult. homolog 3 (E. coli) 1.25
T-cell leukemia/lymphoma 6 1.25
tumor protein D52-like 1 1.25
KIAA0602 protein 1.25
likely ortholog of maternal embryonic leucine zipper kinase 1.25
tubby like protein 2 1.25
a disintegrin and metalloproteinase domain 29 1.25
fragile X mental retardation, autosomal homolog 1 1.25
kynureninase (L-kynurenine hydrolase) 1.25
phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila) 1.25
hypothetical protein DKFZ547M236 1.25
Consensus includes gb:AL050043.1 /DEF=Homo sapiens mRNA; cDNA DKFZ566M0524 (from clone DKFZ566M0524). /FEA=mRNA /DB_XREF=gi:484285 /UG=Hs.3c /AL050043 1.25
gamma-glutamyltransferase 1 1.25
hypothetical protein FLJ23186 1.25
chromosome 21 open reading frame 108 1.25
cone-rod homeobox 1.25
integrin, beta 4 1.25
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1) 1.25
transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma) 1.25
potassium inwardly-rectifying channel, subfamily J, member 1 1.25
plasminogen activator, urokinase 1.25
apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2 1.25
leukocyte cell-derived chemotaxin 2 1.25
Consensus includes gb:AL037069 /FEA=EST /DB_XREF=gi:4073996 /DB_XREF=est:q82h04.x1 /CLONE=IMAGE:2009047 /UG=Hs.17518 Homo sapiens cigs mRNA, par A1337069 1.25
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 2 1.24
parathyroid hormone-like hormone 1.24
aniraxin A3 1.24
hypothetical protein FLJ10998 1.24
prostaglandin E receptor 4 (subtype EP4) 1.24
myosin VA (heavy polypeptide 12, myosin) 1.24
cerebellin 1 precursor 1.24
Consensus includes gb:BE223030 /FEA=EST /DB_XREF=gi:5910348 /DB_XREF=est:hu46c02.x1 /CLONE=IMAGE:3173090 /UG=Hs.301281 Homo sapiens mRNA, chrotr BE223030 1.24
hydroxyacid oxidase 2 (long chain) 1.24
chromosome 8 open reading frame 1 1.24
VPS10 domain receptor protein 1.24
Consensus includes gb:AK024944.1 /DEF=Homo sapiens cDNA: FLJ21291 fts, clone COL01963. /FEA=mRNA /DB_XREF=gi:10437366 /UG=Hs.287657 Homo sapiens cD AK024944 1.24
LIM homeobox protein 1 1.24
soluble carrier family 35 (UDP-galactose transporter), member 2 1.24
aristaeless-like homeobox 3 1.24
hypothetical protein PR02214 1.24
lymphocyte antigen 117 1.24
carbonic anhydrase XIV 1.24
protein kinase, cAMP-dependent, catalytic, alpha 1.24
dishevelled, dish homolog 2 (Drosophila) 1.24
GTPase regulator associated with the focal adhesion kinase pp125 1.24
Consensus includes gb:AB051438.1 /DEF=Homo sapiens mRNA for KIAA1651 protein, partial cds. /FEA=mRNA /GEN=KIAA1651 /PROD=KIAA1651 protein /DB_XREF=gi:AB051438 1.24

organic cation transporter OKB1	AL050350	1.24
hook 1 protein	NM_015888	1.24
hypothetical protein FLJ20555	NM_017877	1.24
guanine nucleotide binding protein (G protein) alpha 12	AK024696	1.24
Consensus includes gb:AL133576.1/DEE=Homo sapiens mRNA; cDNA DKFZp434L0650 (from clone DKFZp434L0650); partial cds. /FEA=mrRNA /GEN=DKFZp434L0650	AL133576	1.24
Consensus includes gb:BF447105 /FEA=EST /DB_XREF=est/749607.x1 /CLONE=IMAGE3649189 /UG=Hs.281706 sordilin 1	BF447105	1.24
Fc fragment of IgG, low affinity Iib, receptor for (CD32)	M31933	1.24
Consensus includes gb:AV733950 /FEA=EST /DB_XREF=gi:10851495 /DB_XREF=est/AV733950 /CLONE=cdAAADG12 /UG=Hs.326035 early growth response 1 /FL=gb:M	AV733950	1.24
TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 250 kD	M73444	1.24
ribosomal protein S8 kinase, 90kD, polypeptide 5	AF074393	1.24
v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)	NM_005465	1.24
cisplatin resistance associated	NM_006687	1.24
caldesmon 1 (fast-twitch, skeletal muscle)	NM_001231	1.24
ubiquitin-conjugating enzyme E2B (RAD8 homolog)	BC005979	1.24
glucosidase, alpha: acid (Pompe disease, glycogen storage disease type II)	NM_000152	1.24
KIAA1128 protein	AI478879	1.24
hypothetical protein PRO1163	NM_018576	1.24
hypothetical protein MGC5487	BC001305	1.24
myxovirus (influenza) resistance 2, homolog of murine	NM_002463	1.24
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 1	NM_014512	1.24
ATP synthase, H ⁺ -transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	D13119	1.24
activin A receptor, type IB	NM_020327	1.24
inactive progesterone receptor, 23 kD	BE903880	1.24
phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 duncce homolog, Drosophila)	U83712	1.24
hypothetical protein FLJ14249 similar to HS1 binding protein 3	NM_022460	1.24
myotubularin related protein 7	AF073482	1.24
R4335, member RAS oncogene family	NM_008861	1.24
nuclear receptor coactivator 2	AW027474	1.24
fibrinogen, A alpha polypeptide	NM_000508	1.24
cell death-inducing DFFA-like effector b	NM_014430	1.24
nephtrin related gene	NM_018240	1.24
unnamed HERV-H protein	U88896	1.24
phospholipase A2, group IVC (cytosolic, calcium-independent)	AF065214	1.24
beta-site APP-cleaving enzyme 2	NM_012105	1.24
endoplasmic reticulum resident protein 58; hypothetical protein MGC5302	NM_024069	1.24
cysteine and glycine-rich protein 3 (cardiac LIM protein)	NM_003476	1.24
t-complex 1	BF224073	1.24
cystic fibrosis transmembrane conductance regulator, ATP-binding cassette (sub-family C, member 7)	W60595	1.24
peroxisome proliferative activated receptor, gamma, coactivator 1	NM_013261	1.24
connexin46.6	BF061637	1.24
glutamate receptor, metabotropic 6	NM_000843	1.24
glycoprotein (transmembrane) nmb	NM_002510	1.24
glutathione S-transferase theta 1	NM_000853	1.24
small inducible cytokine subfamily B (Cys-X-Cys), member 14 (BR4K)	NM_004887	1.24
mitogen-activated protein kinase kinase kinase 12	BE222201	1.24
chromosome 8 open reading frame 1	BF057458	1.24
tripartite motif-containing 28	BE967532	1.24
UDP-Gal-betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 3	AB050656	1.24
ribosomal protein L5	AL137858	1.24
B-cell CLL/lymphoma 9	NM_004326	1.24
oculocerebrorenal syndrome of Lowe	NM_000276	1.24
ATPase, H ⁺ /K ⁺ exchanging, alpha polypeptide	NM_000704	1.24
gamma-glutamyltransferase 1	L20492	1.24
hepatocyte nuclear factor 3, beta	AI693985	1.24
oxytocin receptor, family 2, subfamily B, member 2	AJ302584	1.24
erythrocyte membrane protein band 7.2 (stomatol)	AF130103	1.24
nuclear matrix protein p84	NM_005131	1.24
hypothetical protein from clone 643	BC003379	1.24
calcium/calmodulin-dependent protein kinase (CaM kinase) II beta	AF078803	1.24
BCL2-related protein A1	NM_004048	1.24
hypothetical protein FLJ13892	NM_025017	1.24
pervalbumin	NM_002854	1.24
sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A	NM_020796	1.24

thyroid hormone receptor interactor 10	NM_004240	1.24
hypothetical protein MGC10327	TS4159	1.24
spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal dominant, ataxin 2)	NM_002973	1.24
leukocyte receptor cluster (LRC) member 4	NM_024298	1.24
deoxycycline kinase	NM_000788	1.24
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homologous to yeast nitrogen permease (candidate tumor suppressor)	NM_002177	1.24
interferon, omega 1	NM_004072	1.24
chemokine-like receptor 1	NM_013959	1.24
neuregulin 1	AA053830	1.24
C-terminal binding protein 1	NM_006903	1.24
inorganic pyrophosphatase	NM_014822	1.24
loss of heterozygosity, 11, chromosomal region 2, gene A	AI753638	1.24
orphaned binding protein-like 2	AL049925	1.24
DKFZP547G0910 protein	BE973687	1.24
Consensus includes gb:AW975117 /FEA=EST /DB_XREF=gi:8166322 /DB_XREF=est:EST387223 /UG=Hs.145501 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAM1 AW975117	NM_018067	1.24
hairy homolog (Drosophila)	NM_018235	1.24
hypothetical protein FLJ10350	NM_024939	1.24
hypothetical protein FLJ10830	BC005153	1.24
hypothetical protein FLJ21918	NM_000760	1.24
reghilin 3A-like (without C2 domains)	X87870	1.24
cytochrome P450, subfamily VIIA (cholesterol 7 alpha-monooxygenase), polypeptide 1	NM_000512	1.24
hepatocyte nuclear factor 4, alpha	AJ300481	1.24
galactosamine (N-acetyl)-6-sulfate sulfatase (Morquio syndrome, mucopolysaccharidosis type IVA)	AF072164	1.24
chromosome 11 open reading frame 25	NM_005074	1.24
HFSE-1 protein	NM_001195	1.24
solute carrier family 17 (sodium phosphate), member 1	NM_001502	1.24
beaded filament structural protein 1, flensin	NM_005069	1.24
single-minded homolog 2 (Drosophila)	NM_001559	1.24
glycoprotein 2 (zymogen granule membrane)	NM_023067	1.24
interleukin 12 receptor, beta 2	NM_006464	1.24
forkhead box L2	BC006454	1.24
trans-golgi network protein 2	AK025116	1.24
growth arrest-specific 7	NM_002870	1.24
hypothetical protein FLJ21483	NM_016116	1.24
RAB13, member RAS oncogene family	NM_014448	1.24
ankyrin repeat and SOCS box-containing 4	NM_006561	1.24
Rho guanine exchange factor (GEF) 16	AF118663	1.24
CUG triplet repeat, RNA binding protein 2	AF118663	1.24
maternally expressed 3	AF118663	1.24
Consensus includes gb:AF035307.1 /DEF=Homo sapiens clone 23785 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2681068 /UG=Hs.184697 Homo sapiens clone 23785 AF035307	AF110314	1.24
Consensus includes gb:AL117549.1 /DEF=Homo sapiens mRNA: cDNA DKFZp566H0123 (from clone DKFZp566H0123). /FEA=mRNA /DB_XREF=gi:5912083 /UG=Hs.26 AL117549	AF172331	1.24
poliovirus receptor-related 1 (herpesvirus entry mediator C; nectin)	NM_003489	1.24
regenerating islet-derived 1 alpha (pancreatic stone protein, pancreatic thread protein)	AC002398	1.24
nuclear receptor interacting protein 1	AB002360	1.24
hypothetical protein F25965	BE138825	1.24
MCF-2 cell line derived transforming sequence-like	AI571208	1.24
major histocompatibility complex, class I, F	NM_014095	1.24
C18B11 homolog (44.9kD)	M88013	1.24
PRO1600 protein	NM_002837	1.24
guanine nucleotide binding protein (G protein), alpha 11 (Gq class)	AF113663	1.24
protein tyrosine phosphatase, receptor type, B	NM_030896	1.24
Consensus includes gb:AF113683.1 /DEF=Homo sapiens clone FLB3521. /FEA=mRNA /DB_XREF=gi:6855610 /UG=Hs.306569 Homo sapiens clone FLB3521	W28800	1.24
Consensus includes gb:AF141901.1 /DEF=Homo sapiens GKAPSAPAP interacting protein (SHANK) mRNA, partial cds. /FEA=mRNA /GEN=SHANK /PROD=GKAPSAPAP AF141901	BE856374	1.24
Consensus includes gb:W28800 /FEA=EST /DB_XREF=est:52g12 /UG=Hs.169802 ESTs	NM_024005	1.24
hypothetical protein FLJ13520	NM_017566	1.24
Consensus includes gb:W28800 /FEA=EST /DB_XREF=est:52g12 /UG=Hs.169802 ESTs	AL136460	1.24
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 3	NM_016079	1.24
hypothetical protein FLJ12552	AW474615	1.24
chromosome 9 open reading frame 7	NM_013939	1.24
proliferation-associated 204, 30kD		
CGI-149 protein		
Consensus includes gb:AW474615 /FEA=EST /DB_XREF=gi:7044721 /DB_XREF=est:xy18602.x1 /CLONE=IMAGE:2653506 /UG=Hs.250988 ESTs		
olfactory receptor, family 10, subfamily H, member 2		

sialidase 3 (membrane sialidase)	NM_006656	1,24
glycerol kinase	AA292874	1,24
KIAA0060 protein	AA628780	1,24
potassium inwardly-rectifying channel, subfamily J, member 8	BF514158	1,24
MBIP protein	NM_016586	1,24
doublecortin and Cam kinase-like 1	NM_004734	1,24
leucine zipper-EF-hand containing transmembrane protein 1	NM_012318	1,24
histone deacetylase 8	NM_014707	1,24
Ca11	NM_019617	1,24
squamous cell carcinoma antigen recognized by T cell	NM_013352	1,24
interferon induced transmembrane protein 3 (1-8U)	BF338947	1,24
tubulin-specific chaperone d	AW247323	1,24
hypothetical protein FLJ13954	NM_025020	1,24
tyr-related kinase	NM_002031	1,24
putative GTP-binding protein similar to RAB11C	AL037167	1,24
KIAA0053 gene product	D29642	1,24
hypothetical protein FLJ10565	NM_018140	1,24
mitochondrial ribosomal protein S19C	NM_016087	1,24
KIAA1193 protein	AW003889	1,24
lipopolysaccharide binding protein	NM_004139	1,24
KIAA1048 protein	NM_014911	1,24
carbamoyl-phosphate synthetase 1, mitochondrial	W80357	1,24
patched related protein translocated in renal cancer	AF064801	1,24
glycine receptor, beta	NM_000824	1,24
protein tyrosine phosphatase, non-receptor type 3	NM_002829	1,24
erythroid differentiation-related factor	NM_016633	1,24
actin-like 6	NM_016188	1,24
Consensus includes gb:BE515346 /FEA=EST /DB_XREF=gi:9722561 /DB_XREF=est:601235986F1 /CLONE=IMAGE:3608205 /UG=Hs.278871 ESTs, Weakly similar to S	BE515346	1,24
RAB3B, member RAS oncogene family	BC005035	1,24
ribosomal protein S28	BF431363	1,24
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phosphoglycerate mutase 2 (muscle)	NM_000290	1,24
nitric oxide synthase 1 (neuronal)	NM_006620	1,24
intersectin 2	AF182199	1,24
angiopoietin-like 2	A074333	1,24
hypothetical protein FLJ12789	NM_022495	1,24
homeo box (expressed in ES cells) 1	U82811	1,24
epithelial membrane protein 1	NM_001423	1,24
Consensus includes gb:AL49301.1 /DEF=Homo sapiens mRNA: cDNA DKFp564P073 (from clone DKFp564P073). /FEA=mRNA /DB_XREF=gi:4500087 /UG=Hs.2785	AL49301	1,24
mucin 1, transmembrane	A010889	1,24
hypothetical protein FLJ23604	NM_025084	1,24
upstream binding protein 1 (LBP-1a)	NM_014517	1,24
MCF-2 cell line derived transforming sequence-like	AB002360	1,24
DNA segment, single copy probe LNS-CAI/LNS-CAII (deleted in polyposis	BC000232	1,24
Wiskott-Aldrich syndrome-like	BE504979	1,24
short-chain alcohol dehydrogenase family member	NM_005794	1,24
chorea acanthocytosis	BF573874	1,24
KIAA0828 protein	AK025372	1,24
fibroblast growth factor 2 (basic)	M27968	1,24
Consensus includes gb:AA156777 /FEA=EST /DB_XREF=gi:1728392 /DB_XREF=est:z18c08.s1 /CLONE=IMAGE:502286 /UG=Hs.25431 KIAA1219 protein	AA156777	1,24
lysosomal serylase-like protein 1	NM_020354	1,24
chromosome condensation protein G	NM_022346	1,24
S100-type calcium binding protein A14	NM_020672	1,24
potassium inwardly-rectifying channel, subfamily J, member 16	NM_018656	1,24
telomerase reverse transcriptase	NM_003219	1,24
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 28	A082937	1,24
hypothetical protein FLJ10743	NM_018201	1,24
interleukin 26	NM_018402	1,24
connector enhancer of KSR-like (Drosophila kinase suppressor of ras)	NM_008314	1,24
B1 for mucin	X83412	1,24
hypothetical protein	BF572868	1,24
RNA guanylyltransferase and 5'-phosphatase	NM_003800	1,24
aspartyl aminopeptidase	A050563	1,24

[illegible]

CGI-04 protein 1.23
homoeo box A2 1.23
potassium channel, subfamily K, member 5 (TASK-2) 1.23
hypothetical protein FLJ11125 1.23
SMT3 suppressor of mit two 3 homolog 2 (yeast) 1.23
hypothetical protein FLJ13231 1.23
wingless-type MMTV integration site family, member 11 1.23
gap junction protein, alpha 7, 45KD (connexin 45) 1.23
KIAA0090 protein 1.23
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Consensus includes gb:AA401963 /FEA=EST /DB_XREF=gi:2055965 /DB_XREF=IMAGE:741710 /UG=Hs.194107 ESTs. Moderately similar to AI AA401963 1.23
mesothelin 1.23
Consensus includes gb:A1927479 /FEA=EST /DB_XREF=gi:5663443 /DB_XREF=IMAGE:2462603 /UG=Hs.165590 ribosomal protein S13 1.23
EBIN1 binding protein 2 1.23
mitogen-activated protein kinase 9 1.23
T-box 5 1.23
Consensus includes gb:BE783279 /FEA=EST /DB_XREF=gi:10204477 /DB_XREF=est601470835F1 /CLONE=IMAGE:3873801 /UG=Hs.33862 ESTs 1.23
BTB and CNC homology 1, basic leucine zipper transcription factor 1 1.23
arradillo repeat gene deletion in velocardiofacial syndrome 1.23
seitin/SUMO-specific protease 3 1.23
Consensus includes gb:AL043266 /FEA=EST /DB_XREF=gi:5935844 /DB_XREF=estDKFZp434L1423_s1 /CLONE=DKFZp434L1423 /UG=Hs.111334 ferritin, light polypeptide AL043266 1.23
gb:BC002778.1 /DEF=Homo sapiens. Similar to myosin light chain 2, precursor lymphocyte-specific, clone MGC:3479, mRNA, complete cds. /FEA=mRNA /PROD=Similar BC002778 1.23
solute carrier family 8 (neurotransmitter transporter, GABA), member 11 1.23
transcription factor binding to IGHM enhancer 3 1.23
methionine sulfoxide reductase A 1.23
MRS2-like, magnesium homeostasis factor (S. cerevisiae) 1.23
H1 histone family, member T (testis-specific) 1.23
Consensus includes gb:AI537540 /FEA=EST /DB_XREF=gi:4451675 /DB_XREF=esttp04g11.x1 /CLONE=IMAGE:2188852 /UG=Hs.71168 Homo sapiens clone 24674 mR AI537540 1.23
Consensus includes gb:A1979278 /FEA=EST /DB_XREF=gi:5604295 /DB_XREF=IMAGE:2515854 /UG=Hs.177482 ESTs. Weakly similar to KIAA A1979278 1.23
KIAA0321 protein 1.23
somatostatin 1.23
glycoprotein V (platelet) 1.23
CTAGE-1 protein 1.23
vasoactive intestinal peptide receptor 1 1.23
putative homeodomain transcription factor 1 1.23
KIAA0285 gene product 1.23
G protein-coupled receptor 12 1.23
putative N-acetyltransferase Camello 2 1.23
Consensus includes gb:AA045174 /FEA=EST /DB_XREF=gi:1523376 /DB_XREF=est:166a07.s1 /CLONE=IMAGE:487764 /UG=Hs.16163 Homo sapiens mRNA; cDNA DI AV727634 1.23
fatty-acyl-Coenzyme A ligase, long-chain 6 1.23
Rho guanine exchange factor (GEF) 15 1.23
transglutaminase 4 (prostate) 1.23
Consensus includes gb:AL049337.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564P016 (from clone DKFZp564P016). /FEA=mRNA /DB_XREF=gi:4500118 /UG=Hs.1325 AL049337 1.23
N-acetylglucosamine 6-phosphatase (acid ceramidase) 1.23
translocase of outer mitochondrial membrane 34 1.23
kalikrein 8 (neuropilin/ovastin) 1.23
ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9 1.23
serine protease inhibitor, Kazal type 4 1.23
XO4603 B subunits thnC, thnB genes corresponding to nucleotides 248-2229 of XO4603 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively) 1.23
trefoil factor 3 (intestinal) 1.23
phospholipase A2, group IID 1.23
fructosyltransferase 7 1.23
Consensus includes gb:AL136548.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G18121 (from clone DKFZp761G18121); complete cds. /FEA=mRNA /GEN=DKFZp761 AL136548 1.23
serum/glucoconoid regulated kinase 2 1.23
dysfem intermediate chain 2 1.23
Consensus includes gb:AL080088.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J062 (from clone DKFZp564J062). /FEA=mRNA /DB_XREF=gi:5262475 /UG=Hs.27253 AL080088 1.23
topoisomerase (DNA) III alpha 1.23
Consensus includes gb:W87634 /FEA=EST /DB_XREF=gi:1401698 /DB_XREF=est:zh87b09.s1 /CLONE=IMAGE:417113 /UG=Hs.17947 ESTs. Weakly similar to K02F3.11 W87634 1.23
Jagged 2 1.23
hypothetical protein MGC2376 1.23
CGI-79 protein 1.23

secreted phosphoprotein 1 (osteopontin, bone sialoprotein 1, early T-lymphocyte activation 1) 1.23
 tyrosinase-related protein 1 1.23
 lymphocyte antigen 117 1.23
 myosin 1A 1.23
 lymphocyte antigen 117 1.23
 pleckstrin B2 1.23
 Cluster Ind. T87245:y91b02.r1 Homo sapiens cDNA, 5' and 3' ends IMAGE:115583 /clone_end=5 /gb=T87245 /gi=715597 /ug=Hs.16336 /len=412 1.23
 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 1 1.23
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 4 1.23
 CD84 antigen (leukocyte antigen) 1.23
 hypothetical protein FLJ20837 1.23
 Consensus includes gb:AK024949.1 /DEF=Homo sapiens cDNA: FLJ21296 /ts, clone COL020293 /FEA=mRNA /DB_XREF=gi:10437372 /UG=Hs.306734 Homo sapiens cDNA: AK024949 1.23
 calcium binding protein 1 (calbindin) 1.23
 organic carbon transporter 1.23
 tumor necrosis factor receptor superfamily, member 21 1.23
 prostate differentiation factor 1.23
 kinesin family member 3A 1.23
 TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31 kD 1.23
 Consensus includes gb:A081543 /FEA=EST /DB_XREF=gi:3416335 /DB_XREF=est:cn04g12.x1 /CLONE=IMAGE:1555750 /UG=Hs.300851 ESTs, Highly similar to T0870 1.23
 B-factor, propeptide 1.23
 pavin, beta 1.23
 midline 2 1.23
 translocase of inner mitochondrial membrane 10 homolog (yeast) 1.23
 hypothetical protein FLJ12190 1.23
 Consensus includes gb:AK025038.1 /DEF=Homo sapiens cDNA: FLJ21385 /ts, clone COL03356 /FEA=mRNA /DB_XREF=gi:10437474 /UG=Hs.306750 Homo sapiens cDNA: AK025038 1.23
 Consensus includes gb:AL038824 /FEA=EST /DB_XREF=gi:5407955 /DB_XREF=est:DKFp56602446_s1 /CLONE=DKFp56602446 /UG=Hs.12840 Homo sapiens germ 1.23
 forkhead box O3A 1.23
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 6 1.23
 granzyme M (lymphocyte met-ase 1) 1.23
 neurotrophic tyrosine kinase, receptor, type 3 1.23
 death effector filament-forming Ced-4-like apoptosis protein 1.23
 likely ortholog of mouse p16i like homolog 1 (Orsophila)-like 1.23
 c-fos induced growth factor (vascular endothelial growth factor D) 1.23
 gonadotropin-releasing hormone receptor 1.23
 ubiquitin specific protease 19 1.23
 Consensus includes gb:A0478795 /FEA=EST /DB_XREF=gi:4373608 /DB_XREF=est:tm5207.x1 /CLONE=IMAGE:2161741 /UG=Hs.41123 ESTs, Moderately similar to T1 1.23
 Consensus includes gb:A346431 /FEA=EST /DB_XREF=gi:4083637 /DB_XREF=est:qp50g09.x1 /CLONE=IMAGE:1926496 /UG=Hs.75188 wee1+ (S. pombe) homolog 1.23
 phosphatidylinositol 4-kinase type II 1.23
 hypothetical protein FLJ20618 1.23
 phospholipid scramblase 1 1.23
 Consensus includes gb:AW974666 /FEA=EST /DB_XREF=gi:8165852 /DB_XREF=est:EST386755 /UG=Hs.293024 ESTs 1.23
 v-abl Abelson murine leukemia viral oncogene homolog 2 (arg, Abelson-related gene) 1.23
 mitogen inducible 2 1.23
 MAX protein 1.23
 hypothetical protein from EUROMAGE 511235 1.23
 EGF-containing fibulin-like extracellular matrix protein 1 1.23
 T-box 1 1.23
 KIAA0590 gene product 1.23
 Consensus includes gb:AW971248 /FEA=EST /DB_XREF=gi:8161093 /DB_XREF=est:EST383337 /UG=Hs.291289 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAM1 1.23
 chondroitin sulfate proteoglycan 5 (neuroglycan C) 1.23
 titin-cap (telethonin) 1.23
 Mas4 homolog 1.23
 adaptor-related protein complex 3, delta 1 subunit 1.23
 fatty acid binding protein 5 (psoriasis-associated) 1.23
 KIAA1046 protein 1.23
 hypothetical protein FLJ10534 1.23
 KIAA0140 gene product 1.23
 collagen, type V, alpha 3 1.23
 nuclear receptor subfamily 0, group B, member 1 1.23
 hypothetical protein MGC4655 1.23
 atrophin-1 interacting protein 1; activin receptor interacting protein 1 1.23
 transcription elongation factor B (SIII), polypeptide 2 (18kD, elongin B) 1.23
 hypothetical protein MGC5590 1.23

Rhesus blood group-associated glycoprotein
Down syndrome critical region gene 1
gb:BC000651.1 /DEF=Homo sapiens, Similar to solute carrier family 1 (glutamate transporter), member 7, clone MGC:2078, mRNA, complete cds. /FEA=mRNA /PROD=SI
HX44 antigen
Consensus includes gb:AA576497 /FEA=EST /DB_XREF=gi:2353597 /DB_XREF=est:nm:7607 s1 /CLONE=IMAGE:1074133 /UG=Hs.252497 ESTs
hypothetical protein FLJ22839
nuclear factor of activated T-cells 5, tonically-responsive
protein kinase, cAMP-dependent, regulatory, type I, beta
KIAA0740 gene product
xylosylprotein beta1.4-galactosyltransferase, polypeptide 7 (galactosyltransferase I)
KIAA0887 protein
Consensus includes gb:D60132 /FEA=EST /DB_XREF=gi:961771 /DB_XREF=est:HUM087G05A /CLONE=GEN-087G05 /UG=Hs.79581 Human clone 23560 mRNA seque
D60132
aromatase-derived factor 2
NM_006923
non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase)
BCL2-associated athanogene 3
NM_003551
integral membrane protein 2A
NM_004281
cortactin SH3 domain-binding protein
NM_004867
sialic acid binding Ig-like lectin 7
BF435773
solute carrier family 4, anion exchanger, member 3
NM_016543
ATPase, Ca++ transporting, plasma membrane 3
NM_005070
tumor necrosis factor receptor superfamily, member 10d, decoy with truncated death domain
AF021233
monokine induced by gamma interferon
NM_002416
very long-chain acyl-CoA synthetase homolog 2
NM_012254
Charol-Leyden crystal protein
NM_001828
myeloid/lymphoid or mixed-lineage leukemia (t(11)thorax homolog, Drosophila)
AA704766
filamin-like 1
NM_004467
protein tyrosine phosphatase, receptor type, E
AA775177
PBX/Ino1d 1 homeobox 2
AK032782
pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1)
BC005918
ATPase, Ca++ transporting, cardiac muscle, fast twitch 1
NM_004320
Consensus includes gb:BE465371 /FEA=EST /DB_XREF=gi:5511146 /DB_XREF=est:hw15c12.x1 /CLONE=IMAGE:3182998 /UG=Hs.227170 ESTs
par-6 partitioning defective 6 homolog beta (C. elegans)
NM_002218
inter-alpha (globulin) inhibitor H4 (plasma kallikrein-sensitive glycoprotein)
BC005686
putative selenocysteine lyase
NM_016558
irradiation-induced protein 4
NM_003979
heterogeneous nuclear ribonucleoprotein A1
AW450929
retinoic acid induced 3
NM_016346
chromosome 5 open reading frame 4
NM_000898
monomine oxidase B
NM_002581
pregnancy-associated plasma protein A
NM_002742
protein kinase C, mu
NM_006681
neuromedin U
NM_005584
KIAA0009 gene product
A006835
mab-21-like 1 (C. elegans)
AA761181
RNA, U17D small nucleolar
AC004808
CD24 antigen (small cell lung carcinoma cluster 4 antigen)
NM_006059
olfactory receptor, family 4, subfamily F, member 3
NM_018495
lamtinh, gamma 3
gb:NM_018495.3 /DEF=Homo sapiens NAG22 protein (LOC55873), mRNA. /FEA=mRNA /GEN=LOC55873 /PROD=NAG22 protein /DB_XREF=gi:13236500 /UG=Hs.2830
schwannomin interacting protein 1
NM_014575
KIAA0883 protein
NM_014954
Consensus includes gb:AK023515.1 /DEF=Homo sapiens cDNA FLJ13453 fts, clone PLACE1003205. /FEA=mRNA /DB_XREF=gi:10435471 /UG=Hs.301715 Homo sapier
AK023515
Kruppel-like factor 12
NM_016562
hypothetical protein PRO1483
uroplakin 1B
NM_016285
hypothetical protein PRO2176
phosphatidylinositol glycan, class A (paroxysmal nocturnal hemoglobinuria)
NM_018515
guanine nucleotide binding protein (G protein), alpha 14
NM_002641
sex hormone-binding globulin
NM_004297
Consensus includes gb:AL137547.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B1120 (from clone DKFZp434B1120). /FEA=mRNA /DB_XREF=gi:6808230 /UG=Hs.25 AL137547
ATP-binding cassette, sub-family C (CFTR/MRP), member 8
BC000122
hypothetical protein FLJ00001
L78207
asialoglycoprotein receptor 2
NM_001181

phosphomannomutase 2	NM_000303	1.23
Rat guanine nucleotide exchange factor RagGSP1A	NM_014636	1.23
cytoplasmic FMRP interacting protein 1	BC005097	1.23
Consensus includes gb:A1351043 /FEA=EST /DB_XREF=est:q22a08.x1 /CLONE=IMAGE:1948310 /UG=Hs.279477 ESTs	A1351043	1.23
KIAA0913 protein	W69120	1.23
hypothetical protein FLB3342	NM_020118	1.23
active carrier family 17 (sodium phosphate), member 2	NM_005835	1.23
DNA segment on chromosome 12 (unique) 2469 expressed sequence	NM_007360	1.23
calreticulin	AD000992	1.23
adenosine A2b receptor	NM_000676	1.23
mammaglobin 1	NM_002411	1.23
SP110 nuclear body protein	NM_004510	1.23
myeloproliferative leukemia virus oncogene	NM_005373	1.23
cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila)	D87469	1.23
hypothetical protein FLJ11539	NM_024748	1.23
fibroblast growth factor 18	NM_003862	1.23
immunoglobulin heavy constant mu	X95660	1.23
Consensus includes gb:AA654586 /FEA=EST /DB_XREF=gi:2590740 /DB_XREF=est:n0411.s1 /CLONE=IMAGE:1192173 /UG=Hs.325505 ESTs, Moderately similar to H	AA654586	1.23
a disintegrin and metalloproteinase domain 8	A814527	1.22
caveolin 2	BF187655	1.22
teratocarcinoma-derived growth factor 1	NM_003212	1.22
cystatin C (amyloid angiopathy and cerebral hemorrhage)	NM_000099	1.22
src family associated phosphoprotein 1	NM_003726	1.22
gb:AF122828.1 /DEF=Homo sapiens neurofibromatosis type 2 protein isoform Mer151 (NF2) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=NF2 /PROD=	AF122828	1.22
Consensus includes gb:AV088547 /FEA=EST /DB_XREF=gi:6044358 /DB_XREF=est:xd17c08.x1 /CLONE=IMAGE:2594030 /UG=Hs.312076 ESTs, Moderately similar to	AV088547	1.22
BUB1 budding inhibited by benzimidazoles 1 homolog (yeast)	AK023540	1.22
peroxisome receptor 1	NM_000319	1.22
potassium voltage-gated channel, subfamily G, member 1	A332876	1.22
X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound	NM_003399	1.22
immunoglobulin superfamily, member 4	NM_014333	1.22
histamine receptor H3	NM_007232	1.22
Consensus includes gb:AA773554 /FEA=EST /DB_XREF=gi:2825125 /DB_XREF=est:ab61a05.s1 /CLONE=IMAGE:845264 /UG=Hs.234216 Human 40871 mRNA partial s	AA773554	1.22
egrin	AW008051	1.22
thymosin, beta, Identified in neuroblastoma cells	NM_021992	1.22
CD7 antigen (p41)	NM_006137	1.22
rng finger protein (C3HC4 type) 8	NM_003958	1.22
vascular Rab-GAP/TBC-containing	BC001663	1.22
CD38 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II)	NM_005506	1.22
hypothetical protein FLJ20534	NM_017887	1.22
Consensus includes gb:AL356504 /DEF=Human DNA sequence from clone RP1-14N1 on chromosome 1q21.1-21.3 Contains ESTs, GSSs and STSs. Contains the FLG ge	AL356504	1.22
zinc finger protein 181	BG035118	1.22
protease, serine, 8 (proslasin)	NM_002773	1.22
histidine-rich glycoprotein	NM_006412	1.22
KIAA0864 protein	AB020671	1.22
FOXJ2 forkhead factor	NM_018416	1.22
hemochromatosis	BG402460	1.22
calenin (cadherin-associated protein), delta 2 (neural plakophilin-related arm-repeat protein)	U98136	1.22
amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	BC004369	1.22
KIAA1644 protein	BF516072	1.22
hypothetical protein FLJ12969	NM_022838	1.22
Bacteriophage /REF=X03453 /DEF=Bacteriophage P1 cre recombinase protein corresponding to nucleotides 1032-1270 of X03453 /LEN=1058 /5 and -3 represent transcr	X03453	1.22
KIAA0800 gene product	NM_014703	1.22
tissue inhibitor of metalloproteinase 3 (Sorby fundus dystrophy, pseudoinflammatory)	U87195	1.22
Consensus includes gb:AF254357.1 /DEF=Homo sapiens prostate-specific membrane antigen PSM mRNA, exon 8 alternative splice variant, partial cds. /FEA=mRNA /PRI	AF254357	1.22
neurexophilin 4	A1933199	1.22
Consensus includes gb:AV712064 /FEA=EST /DB_XREF=gi:10731370 /DB_XREF=est:AV712064 /CLONE=DCAAUD05 /UG=Hs.9456 SWISNF related, matrix associated,	AV712064	1.22
myomesin 1 (skelemin) (185kD)	NM_003803	1.22
gb-NM_005196.1 /DEF=Homo sapiens centromere protein F (350400KD, miosin) (CENPF), mRNA. /FEA=mRNA /GEN=CENPF /PROD=centromere protein F (350400KD,	NM_005196	1.22
CGB-142	AK001280	1.22
Consensus includes gb:AL031118 /DEF=Human DNA sequence from clone 153G14 on chromosome 6p21.3-22.2. Contains the 3 end of the ZNF164 gene for Kruppel-like	AL031118	1.22
collagen, type IV, alpha 4	D17391	1.22
hypothetical protein FLJ12484	NM_022767	1.22
Consensus includes gb:AF009287.1 /DEF=Homo sapiens clone FBA1 CH-du-chat region mRNA. /FEA=mRNA /DB_XREF=gi:2331069 /UG=Hs.102238 Homo sapiens clone	AF009267	1.22

vascular cell adhesion molecule 1	NM_001078	1.22
displein resistance-associated overexpressed protein	AW089673	1.22
pp28, US snRNP 100 kD protein	AF028402	1.22
Consensus includes gb:AL133089.1 /DEF=Homo sapiens mRNA; cDNA DKFZp34E2028 (from clone DKFZp34E2028) /FEA=mRNA /DB_XREF=gi:5453541 /UG=Hs.27	AL133089	1.22
CUG triplet repeat, RNA binding protein 2	N36839	1.22
run-related transcription factor 2	NM_004348	1.22
ADP-ribosylation factor 4	NM_001660	1.22
likely ortholog of mouse p1w1 like homolog 1 (Drosophila)-like	NM_018068	1.22
SCAN domain-containing 1	AL109965	1.22
major histocompatibility complex, class II, DP alpha 2 (pseudogene)	X01634	1.22
testis-specific kinase 2	NM_007170	1.22
immunoglobulin heavy constant mu	AB035175	1.22
kinase suppressor of ras	AA971768	1.22
DKFZP568N0721 protein	NM_015400	1.22
hypothetical protein FLJ14251	NM_024881	1.22
topoisomerase (DNA) II binding protein	NM_007027	1.22
DKFZP564F0522 protein	AL049843	1.22
hypothetical protein R29124_1	BF110434	1.22
Consensus includes gb:AL353943.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761E0611 (from clone DKFZp761E0611) /FEA=mRNA /GEN=DKFZp761E0611 /PROD=hy	AL353943	1.22
neurochondrin	AB011179	1.22
UL18 binding protein 1	NM_025216	1.22
cytoplasmic linker 2	BC006259	1.22
matrix Gla protein	AL532375	1.22
hypothetical gene CG018	AL049786	1.22
Consensus includes gb:AL307586 /FEA=EST /DB_XREF=gi:4002311 /DB_XREF=est:tb35h12.x1 /CLONE=IMAGE:2056391 /UG=Hs.133130 Homo sapiens mRNA; cDNA	AL307586	1.22
KIAA0417 gene product	AB007877	1.22
Consensus includes gb:AL535683 /FEA=EST /DB_XREF=gi:4449818 /DB_XREF=est:cong1.P11.A9 /UG=Hs.327729 ESTs	AL535683	1.22
ET4-like factor 5 (eta domain transcription factor)	NM_001422	1.22
CD1c antigen, c polypeptide	NM_001785	1.22
bromodomain adjacent to zinc finger domain, 2A	AI853126	1.22
v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian)	NM_001982	1.22
Consensus includes gb:AL109666.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 35907 /FEA=mRNA /DB_XREF=gi:5689800 /UG=Hs.7242 Ho	AL109666	1.22
hypothetical protein MGC12780	BC006312	1.22
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13	AJ001696	1.22
gb:U19346.1 /DEF=Human (tpi-met fusion) oncogene mRNA, complete cds. /FEA=mRNA /GEN=tpi-met fusion /PROD=tpi-met fusion protein /DB_XREF=gi:525085 /FL=gb	U19346	1.22
Consensus includes gb:BF082139 /FEA=EST /DB_XREF=gi:10821049 /DB_XREF=est:7N7210.x1 /CLONE=IMAGE:3481059 /UG=Hs.282387 polymerase (RNA) III (DNA	BF082139	1.22
hypothetical protein DKFZp564D0372	NM_031301	1.22
olfactory receptor, family 2, subfamily H, member 3	NM_007160	1.22
Consensus includes gb:AK022038.1 /DEF=Homo sapiens cDNA FLJ11978 fls, clone HEMBB1001253 /FEA=mRNA /DB_XREF=gi:10433356 /UG=Hs.296679 Homo sapie	AK022038	1.22
hypothetical protein MGC4170	NM_024312	1.22
Notch homolog 2 (Drosophila)	NM_024408	1.22
nuclear receptor subfamily 1, group 1, member 3	NM_005122	1.22
claudin 18	NM_016369	1.22
hypothetical protein	AI005473	1.22
GW128 protein	NM_014052	1.22
T cell receptor beta locus	U86059	1.22
EphA5	X85425	1.22
SRY (sex determining region Y)-box 13	AF116571	1.22
Consensus includes gb:BG109555 /FEA=EST /DB_XREF=gi:12603061 /DB_XREF=est:60228068F1 /CLONE=IMAGE:4367966 /UG=Hs.331207 ESTs	BG109555	1.22
apoptotic protease activating factor	AF248734	1.22
GRB2-related adaptor protein 2	NM_004810	1.22
scellin	NM_003843	1.22
synovial sarcoma, X breakpoint 3	BF184274	1.22
KIAA0514 gene product	NM_014686	1.22
transmembrane protease, serine 2	NM_005656	1.22
Consensus includes gb:AF090886.1 /DEF=Homo sapiens clone HQ0072 /FEA=mRNA /DB_XREF=gi:66890153 /UG=Hs.284192 Homo sapiens clone HQ0072	AF090886	1.22
biothidase	AI787414	1.22
Consensus includes gb:D16471.1 /DEF=Human mRNA, Xq terminal portion. /FEA=mRNA /DB_XREF=gi:495099 /UG=Hs.121571 Human mRNA, Xq terminal portion	D16471	1.22
hypothetical protein FLJ22844	NM_025098	1.22
taste receptor, type 2, member 4	NM_016944	1.22
M10098 Human 18S rRNA gene, complete (L5_M_3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M10098	1.22
solute carrier family 4, anion exchanger, member 2 (erythrocyte membrane protein band 3-like 1)	NM_003040	1.22
gamma-glutamyltransferase-like activity 1	NM_004121	1.22

myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10
 signal-regulatory protein beta 2
 synovial sarcoma, X breakpoint 3
 small inducible cytokine subfamily A (Cys-Cys), member 23
 phospholipase A2-activating protein
 suppressor of Ty 3 homolog (S. cerevisiae)
 small proline-rich protein 2B
 collagen, type I, alpha 2
 tumor protein p53-binding protein
 solute carrier family 4, sodium bicarbonate cotransporter, member 7
 PRO1992 protein
 olfactory receptor, family 1, subfamily A, member 2
 hypothetical protein FLJ23590
 gamma-aminobutyric acid (GABA) A receptor, beta 1
 desmoplakin (DPI, DPII)
 hypothetical protein MGC4694
 carbonic anhydrase XII
 chromosome 5 open reading frame 4
 tubulin, beta polypeptide 4, member Q
 KIAA0853 protein
 Consensus includes gb:AV1715767 /FEA=EST /DB_XREF=gi:10797284 /DB_XREF=est:AV1715767 /CLONE=DCBATH02 /UG=Hs.7780 Homo sapiens mRNA; cDNA DKFZ: AV1715767
 transmembrane 4 superfamily member 1
 hemoglobin, alpha 2
 G protein-coupled receptor C5L2
 nuclear receptor coactivator 1
 potassium voltage-gated channel, KQT-like subfamily, member 4
 similar to rat HREV107
 C3H-type zinc finger protein; similar to D. melanogaster muscledblind B protein
 potassium voltage-gated channel, KQT-like subfamily, member 2
 fucosyltransferase 8 (alpha (1.3) fucosyltransferase)
 Consensus includes gb:AI188346 /FEA=EST /DB_XREF=gi:3739555 /DB_XREF=est:qd08a10.x1 /CLONE=IMAGE:1723098 /UG=Hs.301778 ESTs, Weakly similar to ALU1 AI188346
 aladin 2-binding protein 1
 Consensus includes gb:AV720803 /FEA=EST /DB_XREF=gi:10817955 /DB_XREF=est:AV720803 /CLONE=GLCCMA04 /UG=Hs.293055 ESTs
 DNase II-like acid DNase
 carboxyl ester lipase-like (bile salt-stimulated lipase-like)
 brain-specific membrane-anchored protein
 KIAA0974 protein
 cyclin F
 mannan-binding lectin serine protease 2
 FGFR1 oncogene partner
 Consensus includes gb:BE881590 /FEA=EST /DB_XREF=gi:10330366 /DB_XREF=est:601490008F1 /CLONE=IMAGE:3892465 /UG=Hs.10684 Homo sapiens clone 2442
 sialyltransferase
 reticulon 4
 Consensus includes gb:AA370433 /FEA=EST /DB_XREF=gi:2344433 /DB_XREF=est:m62111.s1 /CLONE=IMAGE:1018125 /UG=Hs.10323 Homo sapiens mRNA from chr
 signal transduction protein (SH3 containing)
 calmodulin-like 3
 KIAA0603 protein
 Consensus includes gb:BG111761 /FEA=EST /DB_XREF=gi:12605267 /DB_XREF=est:602285343F1 /CLONE=IMAGE:4372619 /UG=Hs.8107 Homo sapiens mRNA; cDN
 serum/glucocorticoid regulated kinase 2
 hypothetical protein FLJ12929
 KIAA0492 protein
 Consensus includes gb:AK000257.1 /DEF=Homo sapiens cDNA FLJ20250 fs, clone COLF6635, /FEA=mRNA /DB_XREF=gi:7020217 /UG=Hs.274505 Homo sapiens mRN
 PRO0618 protein
 EGF-like-domain, multiple 6
 H326
 Cyclin M2
 potassium inwardly-rectifying channel, subfamily J, member 10
 hypothetical protein
 type I intermediate filament cytochrome
 fibroblast growth factor 16
 guanylate binding protein 1, interferon-inducible, 67kD
 LIM homeobox protein 2
 cyclic nucleotide gated channel beta 1

245

MHC binding factor, beta
CGI-35 protein
Zinc finger protein 2 heterolog 1 (odd-paired homolog, Drosophila)
melanoma antigen, family A, 11
thyroid stimulating hormone receptor
eyes absent homolog 1 (Drosophila)
Tax interaction protein 1
hypothetical protein FLJ13857
hepclin antimicrobial peptide
KIAA0547 gene product
Consensus includes gb:A198543 /FEA=EST /DB_XREF=gi:3751149 /DB_XREF=est:q49f11.x1 /CLONE=IMAGE:1753389 /UG=Hs.8982 ESTs, Highly similar to KIAA1395 A198543
sarcoglycan, delta (35KD dystrophin-associated glycoprotein)
POZ-75 protein
cytosolic beta-glucosidase
hypothetical protein FLJ22690
H4 histone family, member C
KIAA1219 protein
Consensus includes gb:BG536504 /FEA=EST /DB_XREF=gi:13528050 /DB_XREF=est:602564909F.1 /CLONE=IMAGE:4689314 /UG=Hs.274518 Homo sapiens mRNA; cf BG536504
PBX/noted 1 homeobox 1
meninaglobin 2
FAT tumor suppressor homolog 2 (Drosophila)
chondroitin sulfate proteoglycan BEHAB/brevican
cyclin-dependent kinase inhibitor 1B (p27, Kip1)
Consensus includes gb:BE865029 /FEA=EST /DB_XREF=gi:11768976 /DB_XREF=est:601658812R1 /CLONE=IMAGE:3886131 /UG=Hs.198793 Homo sapiens cDNA: FL BE865029
karyopherin beta 2b, transportin
insulin-like growth factor binding protein 6
solute carrier family 6 (neurotransmitter transporter, creatine), member 8
hypothetical protein AF053356_CDS3
Werner syndrome
Consensus includes gb:AW979018 /FEA=EST /DB_XREF=gi:8170301 /DB_XREF=est:EST391128 /UG=Hs.132670 ESTs
l-mis domain-containing protein
dual specificity phosphatase 4
retinal homeobox protein
hypothetical protein EST00098
tumor necrosis factor (ligand) superfamily, member 11
KIAA0570 gene product
ATP binding protein associated with cell differentiation
chromosome 5 open reading frame 7
secreted protein, acidic, cysteine-rich (osteonectin)
protein tyrosine phosphatase, non-receptor type 1
KIAA1068 protein
wingless-type MMTV integration site family, member 5A
Consensus includes gb:AW404894 /FEA=EST /DB_XREF=gi:6923951 /DB_XREF=est:U1-HF-BLQ-act-f07-0.U1.1 /CLONE=IMAGE:3059844 /UG=Hs.314762 Homo sapiens mRNA; cf AW404894
glucosylase 1
major histocompatibility complex, class II, DQ beta 1
hypothetical protein MGC3035
lipin 2
Consensus includes gb:AW008921 /FEA=EST /DB_XREF=gi:5857699 /DB_XREF=est:wa75a03.x1 /CLONE=IMAGE:2503756 /UG=Hs.13138 Homo sapiens, clone IMAGE AW008921
KIAA0931 protein
Consensus includes gb:AF103295.1 /DEF=Homo sapiens clone N97 immunoglobulin heavy chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin AF103295
KIAA1067 protein
hypothetical protein FLJ10853
CDC42 binding protein kinase beta (DMPK-like)
KIAA0938 protein
LIM and senescent cell antigen-like domains 1
APR-1 protein
guanylate cyclase 1, soluble, alpha 3
annexin A3
cofactor required for Sp1 transcriptional activation, subunit 6 (77KD)
TAF11 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 28 KD
N-methylpurine-DNA glycosylase
chromosome X open reading frame 2
ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)

putative mitochondrial solute carrier	NM_031212	1,21
primitin-like 1 (mouse)	NM_006017	1,21
SRY (sex determining region Y)-box 13	AF083105	1,21
myosin, heavy polypeptide 7, cardiac muscle, beta	A1292276	1,21
nHDC for homolog of Drosophila headcase	NM_016217	1,21
ankyrin 1, erythrocytic	NM_020478	1,21
ankyrin 1, erythrocytic	A1659683	1,21
embryonic ectoderm development	AF099032	1,21
acetylserotonin O-methyltransferase-like	Y15521	1,21
hypothetical protein FLJ20708	NM_017835	1,21
SH3 domain binding glutamic acid-rich protein like	NM_003022	1,21
nebulin	NM_006393	1,21
granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1)	J03189	1,21
oligophrenin 1	NM_002547	1,21
islet cell autoantigen 1 (69kD)	L21181	1,21
hypothetical protein	AL137871	1,21
T cell receptor gamma locus	M30894	1,21
prostaglandin D2 synthase (21kD, brain)	M81900	1,21
hypothetical protein FLJ13287	NM_024647	1,21
hypothetical protein FLJ10094	NM_017893	1,21
src homology three (SH3) and cysteine rich domain	NM_003149	1,21
UDP glycosyltransferase 1 family, polypeptide A3	NM_019093	1,21
proline synthetase co-transcribed (bacterial homolog)	A0021923	1,21
villin 1	NM_007127	1,21
RNA (guanine-7-); methyltransferase	NM_003799	1,21
Consensus includes gb:AL512688.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547J2313 (from clone DKFZp547J2313); /FEA=mRNA /GEN=DKFZp547J2313 /PROD=hyp	AL512688	1,21
zinc finger protein 37 homolog (mouse)	NM_003408	1,21
gb:NM_016184.2 /DEF=Homo sapiens hyaluronic acid receptor (HAR), mRNA, /FEA=mRNA /GEN=HAR /PROD=hyaluronic acid receptor /DB_XREF=agi:10835259 /UG=Hs	NM_016184	1,21
Fancconi anemia, complementation group G	AC004472	1,21
Consensus includes gb:A4558468 /FEA=EST /DB_XREF=gi:2328945 /DB_XREF=estn39606.s1 /CLONE=IMAGE:1015882 /UG=Hs.178665 Homo sapiens immunoglobul	A4558468	1,21
MBD2 (methyl-CpG-binding protein)-interacting zinc finger protein	NM_015517	1,21
interferon, beta 1, fibroblast	NM_002178	1,21
pyruvate dehydrogenase kinase, isoenzyme 2	AL574319	1,21
zinc finger protein 278	AF254087	1,21
inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory)	NM_000362	1,21
hepatocyte growth factor (hepatopoietin A; scatter factor)	U46010	1,21
chloride channel 8	NM_001286	1,21
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1	NM_004727	1,21
Consensus includes gb:AX001442.1 /DEF=Homo sapiens cDNA FLJ10360 fls, clone NT2RP2003533, /FEA=mRNA /DB_XREF=gi:7022701 /UG=Hs.140402 Homo sapiens	AX001442	1,21
Consensus includes gb:AK028273.1 /DEF=Homo sapiens cDNA FLJ22620 fls, clone HSI05629, /FEA=mRNA /DB_XREF=gi:10439077 /UG=Hs.306834 Homo sapiens cOI	AK028273	1,21
killer cell lectin-like receptor subfamily 6, member 1	NM_002258	1,21
hypothetical protein FLJ12988	NM_025148	1,21
solute carrier family 21 (organic anion transporter), member 14	NM_017435	1,21
matrix metalloproteinase 13 (collagenase 3)	NM_002427	1,21
iris-related tyrosine kinase 3 ligand	U03858	1,21
forbeshed box E2	NM_012185	1,21
melastatin 1	N32151	1,21
gonadotropin-releasing hormone receptor	Z81148	1,21
interleukin 10	NM_000572	1,21
KIAA0599 protein	BG223341	1,21
ricized homolog 9 (Drosophila)	NM_003508	1,21
solute carrier family 18 (monocarboxylic acid transporters), member 1	BF511091	1,21
wingless-type Wnt/PCP integration site family, member 4	NM_030761	1,21
heparanase-like protein	NM_021828	1,21
insulin-like 3 (Leydig cell)	A1991894	1,21
hypothetical protein FLJ12983	NM_024656	1,21
interleukin 1 receptor antagonist	BE563442	1,21
hypothetical protein DKFZp762P2111	R61320	1,21
BCE-1 protein	R61320	1,21
heart and neural crest derivatives expressed 1	NM_007005	1,21
coagulation factor XII (Hageman factor)	NM_004821	1,21
KIAA0752 protein	BFS30348	1,21
SET binding factor 1	AF155097	1,21
SET binding factor 1	AU333707	1,21

248

potassium inwardly-rectifying channel, subfamily J, member 5	U39195	1.21	
HSPC063 protein	NM_014155	1.21	
serine protease inhibitor-like, with Kunitz and WAP domains 1 (eppin)	AF286388	1.21	
family with sequence similarity 8, member A1	NM_016255	1.21	
transforming, acidic coiled-coil containing protein 2	NM_006997	1.21	
unt-interacting protein	NM_007178	1.21	
procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2	A1754404	1.21	
transmembrane 4 superfamily member 7	BC000389	1.21	
triple functional domain (TPTRF) interacting	A1797657	1.21	
hypothetical protein FLJ21343	NM_022781	1.21	
small inducible cytokine subfamily C, member 1 (lymphotactin)	U23772	1.21	
solute carrier family 18 (vesicular monoamine), member 1	NM_003053	1.21	
Consensus includes gb:AK024561.1 /DEF=Homo sapiens cDNA: FLJ20908 fts, done ADSE00417, /FEA=mRNA /DB_XREF=gi:10438870 /UG=Hs.306689 Homo sapiens c AK024561	NM_003304	1.21	
transient receptor potential channel 1	NM_003535	1.21	
Consensus includes gb:BC003535.1 /DEF=Homo sapiens, clone IMAGE:3609326, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3609326) /DB_XI BC003535	NM_004106	1.21	
Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide	NM_012124	1.21	
cysteine and histidine-rich domain (CHORD)-containing, zinc binding protein 1	NM_021830	1.21	
chromosome 10 open reading frame 2	NM_002290	1.21	
laminin, alpha 4	NM_030968	1.21	
G protein coupled receptor interacting protein, complement-c1q tumor necrosis factor-related	L27624	1.21	
tissue factor pathway inhibitor 2	U58852	1.21	
nuclear protein, ataxia-telangiectasia locus	AF116713	1.21	
inter-alpha (globulin) inhibitor, H1 polypeptide	A1921320	1.21	
ribosome binding protein 1 homolog 180KD (dog)	BE855983	1.21	
acyl-Coenzyme A carboxylase alpha	AL031073	1.21	
melanoma antigen, family E, 1, cancer/testis specific	NM_000289	1.21	
Consensus includes gb:BC002869.1 /DEF=Homo sapiens, clone IMAGE:3941350, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3941350) /DB_XI BC002869	AB007878	1.21	
KIAA0508 protein	NM_005299	1.21	
G protein-coupled receptor 31	AW024347	1.21	
caspase recruitment domain protein 10	NM_003337	1.21	
ubiquitin-conjugating enzyme E2B (RAD6 homolog)	NM_0167210	1.21	
Consensus includes gb:A1767210 /FEA=EST /DB_XREF=gi:5233719 /DB_XREF=est:tw9405.x1 /CLONE=IMAGE:2400869 /UG=Hs.90449 Human clone 23908 mRNA ser A1767210	D86984	1.21	
dedicator of cyto-kinesis 2	NM_018196	1.21	
epsilon-trimethyllysine hydroxylase	NM_005148	1.21	
unc-119 homolog (C. elegans)	AF098114	1.21	
integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B)	A1671579	1.21	
zinc finger protein 8 (clone HF.18)	NM_022840	1.21	
hypothetical protein FLJ23017	NM_025262	1.21	
GSC protein	AF307080	1.21	
KIAA0768 protein	NM_005203	1.21	
collagen, type XIII, alpha 1	NM_002009	1.21	
fibroblast growth factor 7 (keratinocyte growth factor)	NM_018476	1.21	
brain expressed, X-linked 1	AB051523	1.21	
hypothetical protein FLJ20294	NM_014289	1.21	
calpain 6	U84364	1.21	
glycogenin 2	U58515	1.21	
chitinase 3-like 2	NM_007105	1.21	
solute carrier family 22 (organic cation transporter), member 1-like antisense	NM_024697	1.21	
hypothetical protein FLJ22419	AL022067	1.21	
PR domain containing 1, with ZNF domain	AA524274	1.21	
Splicing factor, arginine/serine-rich, 48KD	R01140	1.21	
heat shock 90KD protein 1, alpha	NM_006998	1.21	
secretagogin	AF254083	1.21	
zinc finger protein 278	A1557467	1.21	
KIAA0478 gene product	U53784	1.21	
paraoxonase 1	NM_025007	1.21	
hypothetical protein FLJ13501	BG398778	1.21	
methylmalonate-semialdehyde dehydrogenase	AU143855	1.21	
KIAA0077 protein	AF211188	1.21	
fibroblast growth factor 18	NM_008731	1.21	
Fukuyama type congenital muscular dystrophy	gb:NM_018407.1 /DEF=Homo sapiens putative integral membrane transporter (LC27), mRNA, /FEA=mRNA /GEN=LC27 /PROD=putative integral membrane transporter /C NM_018407	AA702163	1.21
peanut-like 1 (Drosophila)		1.21	

KIAA0551 protein	N25621	1,21
development and differentiation enhancing factor 2	NM_003887	1,21
G protein-coupled receptor 57	NM_014627	1,21
tachykinin receptor 2	NM_001057	1,21
colony stimulating factor 1 receptor, formerly McDonough feline sarcoma viral (v-fms) oncogene homolog	NM_005211	1,21
cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)	U20498	1,21
Consensus includes gb:M28219.1 (DEF=Homo sapiens low density lipoprotein receptor (FH 10 mutant causing familial hypercholesterolemia) mRNA, 3 end. /FEA=mRNA /M28219		1,21
polymerase (RNA) II (DNA directed) polypeptide D	BF432147	1,21
NAG16 protein	AF210651	1,21
chondroitin 4-O-sulfotransferase 2	NM_018641	1,21
NAO(P)H dehydrogenase, quinone 1	NM_000903	1,21
solute carrier family 6 (neurotransmitter transporter, creatine), member 8	U17986	1,21
solute carrier family 23 (nucleobase transporters), member 1	AF092511	1,21
Protein inhibitor of activated STAT X	AF077954	1,21
platelet/endothelial cell adhesion molecule (CD31 antigen)	AA702701	1,21
KIAA0451 gene product	AA127643	1,21
transferrin (prealbumin, amyloidosis type I)	AF162690	1,21
diaphanous homolog 2 (Drosophila)	NM_008729	1,21
regulator of G-protein signaling 17	NM_012419	1,21
guanylate cyclase 1, soluble, beta 3	W53728	1,21
prostaglandin D2 synthase (21KD, brain)	BC005939	1,21
G protein-coupled receptor 23	NM_005296	1,21
acetylcholinesterase (YT blood group)	NM_015831	1,21
hypothetical protein FLJ12567	NM_024714	1,21
hypothetical protein FLJ13238	NM_024902	1,21
KIAA1624 protein	NM_020960	1,21
keratin, hair, acidic, 8	NM_006771	1,21
capping protein (actin filament) muscle Z-line, beta	BG420053	1,21
acetyl-Coenzyme A carboxylase alpha	AW188201	1,21
ATPase, Class II, type 9B	R50094	1,21
IQ motif containing GTPase activating protein 2	NM_006633	1,21
L38424 B subtilis dapB, jpfF, jpfG genes corresponding to nucleotides 1358-3197 of L38424 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	L38424	1,21
very large G protein-coupled receptor 1	AB014586	1,21
reticulon 1	BC000314	1,21
KIAA1535 protein	BF110802	1,21
peter pan homolog (Drosophila)	AI399889	1,21
KIAA0215 gene product	NM_014735	1,21
glial fibrillary acidic protein	NM_002055	1,21
hypothetical protein FLJ10748	NM_018203	1,21
insulin induced gene 1	BG292233	1,21
ribonuclease, RNase A family, 1 (pancreatic)	NM_002933	1,21
zyxin	BC002323	1,21
Consensus includes gb:BE552409 /FEA=EST /DB_XREF=gi:3794.101 /DB_XREF=est:hy0708.x1 /CLONE=IMAGE:3196647 /UG=Hs.280895 ESTs		1,21
transmembrane protein with EGF-like and two follistatin-like domains 1	BE552409	1,21
3-hydroxy-3-methylglutaryl-Coenzyme A reductase	BF439316	1,21
succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD	NM_000859	1,21
nuclear receptor subfamily 1, group 1, member 2	NM_003001	1,21
neurogenic differentiation 2	NM_003889	1,21
KIAA1111 protein	AB021742	1,21
cell cycle related kinase	AF091081	1,21
KIAA1660 protein	NM_012119	1,21
transmembrane protein 5	NM_014254	1,21
RCD1 required for cell differentiation1 homolog (S. pombe)	AI360850	1,21
mitogen-activated protein kinase 10	NM_002753	1,20
natriuretic peptide receptor A/guanylate cyclase A (atrionatriuretic peptide receptor A)	X15557	1,20
transcription elongation factor B (SII), polypeptide 3 (110KD, elongin A)	NM_003198	1,20
immunoglobulin heavy constant mu	U80164	1,20
pleckstrin and Sec7 domain protein	NM_002779	1,20
histone deacetylase 8	AK024083	1,20
KIAA0672 gene product	NM_014859	1,20
Consensus includes gb:BG032839 /FEA=EST /DB_XREF=gi:12424547 /DB_XREF=est:602300417F1 /CLONE=IMAGE:4401700 /UG=Hs.293341 ESTs, Weakly similar to BG032839		1,20
cytochrome P450, subfamily IIIA (nifedipine oxidase), polypeptide 3	J04449	1,20
zinc finger protein 281	AK025663	1,20

251

Consensus includes gb:BF046750 /FEA=EST /DB_XREF=gi:10763906 /DB_XREF=est:INSSA4 /CLONE=227760 /UG=Hs.274454 ESTs, Weekly similar to 20043994 chr9
 alpha-methylacyl-CoA racemase
 frizzled homolog 2 (Drosophila)
 LIM domain only 6
 BarH-like homeobox 2
 ketch-like 4 (Drosophila)
 Consensus includes gb:AL133269 /DEF=Human DNA sequence from clone RP3-468K3 on chromosome 8q25-26. Contains a beta tubulin pseudogene, STSs and GSSs /F
 olfactory receptor, family 7, subfamily E, member 24 pseudogene
 potassium voltage-gated channel, subfamily H (eag-related), member 4
 RAB3D, member RAS oncogene family
 hypothetical protein
 huntingtin-associated protein 1 (neuroan 1)
 alpha-2-HS-glycoprotein
 fibroblast growth factor 2 (basic)
 complement component 8, alpha polypeptide
 cytochrome P450, subfamily IVA, polypeptide 11
 guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1
 cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9
 likely ortholog of preimplantation protein 3
 centaurin, beta 2
 mitogen-activated protein kinase kinase kinase 5
 KIAA0285 gene product
 hypothetical protein FLJ13511
 I-box and WD-40 domain protein 2
 hypothetical protein DKFZ0781P1010
 WAP four-disulfide core domain 1
 KIAA0431 protein
 Consensus includes gb:AA621286 /FEA=EST /DB_XREF=gi:2523225 /DB_XREF=est:u83e09.s.1 /CLONE=IMAGE:744616 /UG=Hs.179697 Human DNA damage repair a
 hypothetical protein IMPACT
 matrilin 1, cartilage matrix protein
 Consensus includes gb:AK025188.1 /DEF=Homo sapiens cDNA: FLJ21535 fls, clone COL06131 /FEA=mRNA /DB_XREF=gi:10437651 /UG=Hs.306781 Homo sapiens c
 DKFZP564G202 protein
 estrogen receptor binding site associated, antigen, 9
 hypothetical protein similar to beta-transducin family
 cytochrome P450, subfamily IIA, polypeptide 7
 pre-TNK cell associated protein
 KIAA0747 protein
 DKFZP564G092 protein
 Consensus includes gb:U92027.1 /DEF=Human clone 61501 defective mariner transposon Hsmar2 mRNA sequence, /FEA=mRNA /DB_XREF=gi:2052468 /UG=Hs.15921 U92027
 Consensus includes gb:AV760769 /FEA=EST /DB_XREF=gi:10918617 /DB_XREF=est:AV760769 /CLONE=MDSSE09 /UG=Hs.306324 Homo sapiens mRNA; cDNA DKF
 protein tyrosine phosphatase, receptor type, K
 hypothetical protein FLJ20920
 hypothetical protein MGC2479
 KIAA0053 gene product
 myelin oligodendrocyte glycoprotein
 gb:NM_023941.1 /DEF=Homo sapiens hypothetical protein MGC3032 (MGC3032), mRNA, /FEA=mRNA /GEN=MGC3032 /PROD=hypothetical protein MGC3032 /DB_XREF=gi:203941
 GW128 protein
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 1 (anionporter, Na+/H+, amiloride sensitive)
 ATPase, H+-transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1B
 KIAA1405 protein
 lysosomal-associated membrane protein 2
 matorin, ring finger protein, 1
 polymerase (DNA directed), delta 3
 aryl hydrocarbon receptor nuclear translocator
 Consensus includes gb:AL137706.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434A2322 (from clone DKFZp434A2322), /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:203941
 hypothetical protein FLJ20126
 guanine nucleotide binding protein (G protein), alpha 11 (Gq class)
 DNA (cytosine-5-methyltransferase 3-like
 6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 1
 cathepsin B
 Epha7
 RNA binding motif, single stranded interacting protein 1
 Rho GTPase activating protein 1

Consensus includes gb:AB028239 /DEF=Homo sapiens pseudogene for necln /FEA=CDS /DB_XREF=gi:5672690 /UG=Hs.247843 Homo sapiens pseudogene for necln AB028239
 pre-mRNA splicing SR protein RA4 A4648521
 Consensus includes gb:AL049397.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586C1019 (from clone DKFZp586C1019). /FEA=mRNA /DB_XREF=gi:4500188 /UG=Hs.12
 binder of Arf Two BC003087
 KIAA0832 protein AL110250
 potassium voltage-gated channel, shaker-related subfamily, member 1 (episodic ataxia with myokymia) NM_000217
 T cell receptor gamma locus M27331
 KIAA0564 protein BE550798
 KIAA1043 protein NM_028966
 hypothetical protein FLJ22028 NM_024854
 androgen-induced prostate proliferative shut-off associated protein NM_015928
 suppression of tumorigenicity 5 NM_005418
 Consensus includes gb:BG494940 /FEA=EST /DB_XREF=gi:13456455 /DB_XREF=est:602540661F1 /CLONE=IMAGE:4671854 /UG=Hs.554 Sjogren syndrome antigen A BG494940
 Consensus includes gb:BG494940 /FEA=EST /DB_XREF=gi:13456455 /DB_XREF=est:602540661F1 /CLONE=IMAGE:4671854 /UG=Hs.554 Sjogren syndrome antigen A BG494940
 zinc finger protein 214 NM_013248
 integrin, beta 4 NM_000213
 calicum-sensing receptor (hypocalcemic hypercalcemia 1, severe neonatal hyperparathyroidism) U20760
 Consensus includes gb:AK000293 /DEF=Homo sapiens cDNA FLJ20286 fls, clone HEP04358. /FEA=mRNA /DB_XREF=gi:7020278 /UG=Hs.306331 Homo sapiens cDN U20760
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3 U55430
 hypothetical protein FLJ21019 NM_024927
 endolase alpha, lung-specific NM_014355
 Consensus includes gb:A983904 /FEA=EST /DB_XREF=gi:5811123 /DB_XREF=est:w55d05.x1 /CLONE=IMAGE:2561961 /UG=Hs.79732 fibulin 1 A983904
 F-box only protein 24 NM_012172
 hypothetical protein MGC8407 NM_024046
 matrix metalloproteinase 16 (membrane-inserted) U78292
 KIAA0582 gene product NM_014852
 KIAA0329 gene product AB002295
 gb:NM_018069.1 /DEF=Homo sapiens hypothetical protein FLJ10352 (FLJ10352), mRNA. /FEA=mRNA /GEN=FLJ10352 /PROD=hypothetical protein FLJ10352 /DB_XREF=NM_018069
 phosphate cytidylyltransferase 1, choline, alpha isoform NM_005017
 DKFZP564F1422 protein AL049542
 hypothetical protein FLJ21127 NM_024549
 melanoma antigen, family A, 1 (directs expression of antigen MZ2-E) NM_004988
 hydroxyprostaglandin dehydrogenase 15-(NAD) NM_008860
 hypothetical protein FLJ20515 NM_017857
 casein, alpha NM_001890
 golgi autoantigen, golgin subfamily a, 2 L06147
 v-rat tianian leukemia viral oncogene homolog B (ras related; GTP binding protein) BG169873
 splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) BG035151
 hepatic leukemia factor W60800
 Consensus includes gb:AK025097.1 /DEF=Homo sapiens cDNA: FLJ21444 fls, clone COL04437. /FEA=mRNA /DB_XREF=gi:10437544 /UG=Hs.306764 Homo sapiens cC AK025097
 OLF-1/EBF associated zinc finger gene AW149417
 junctional 3 NM_020655
 glycine receptor, alpha 1 (startle disease/hyperkplexia, stiff man syndrome) NM_000171
 bromodomain adjacent to zinc finger domain, 2A NM_000888
 integrin, beta 6 NM_013449
 hypothetical protein FLJ10970 NM_018286
 synovial sarcoma, X breakpoint 3 BF184274
 COX17 homolog, cytochrome c oxidase assembly protein (yeast) NM_005694
 metastasis-associated 1-like 1 AB012922
 C-terminal binding protein 2 AF222711
 retinoic acid receptor, gamma L12060
 hypothetical protein EST00098 NM_022885
 a disintegrin and metalloproteinase domain 28 NM_021777
 cryptochrome 1 (photolyase-like) D63702
 voltage-dependent anion channel 3 BC002456
 Consensus includes gb:AK025247.1 /DEF=Homo sapiens cDNA: FLJ21594 fls, clone COL07053. /FEA=mRNA /DB_XREF=gi:10437718 /UG=Hs.288571 Homo sapiens cC AK025247
 hypothetical protein FLJ22037 NM_024681
 DKFZP564L0862 protein NM_017757
 KIT ligand NM_024087
 interleukin 1 receptor-like 2 NM_000899
 aquaporin 2 (collecting duct) NM_003854
 RNA binding motif protein, Y chromosome, family 1, member A1 NM_000466
 acylserotonin O-methyltransferase-like NM_005058
 1.19 BC002508

chromosome 1 open reading frame 17	AF288389	1.19
ATP-binding cassette, sub-family B (MDR/TAP), member 9	NM_019825	1.19
solute carrier family 25 (mitochondrial carrier, Adlar), member 12	A067457	1.19
DKFZP36682420 protein	BF509857	1.19
insulin receptor substrate 3-like	H07885	1.19
methylmalonate-semialdehyde dehydrogenase	AW612403	1.19
chloride intracellular channel 2	AI768628	1.19
KIAA1418 protein	NM_017783	1.19
Notch homolog 2 (Drosophila)	AF308601	1.19
arylsulfatase A	U62317	1.19
hypothetical protein PRO1048	NM_018497	1.19
thyroid stimulating hormone receptor	BE740743	1.19
tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)	BC002794	1.19
zinc finger protein 142 (clone PHZ-49)	D87073	1.19
KIAA0252 protein	AI92081	1.19
matrix, extracellular phosphoglycoprotein with ASARM motif (bone)	NM_020203	1.19
glycophorin B (includes Ss blood group)	NM_002100	1.19
glycosylphosphatidylinositol specific phospholipase D1	NM_001503	1.19
ZYG homolog	AI991574	1.19
RAR-related orphan receptor C	NM_005060	1.19
adaptor-related protein complex 4, epsilon 1 subunit	NM_007347	1.19
phosphoinositide 3-phosphate-binding protein-2	NM_014935	1.19
homeo box B8	NM_024016	1.19
KIAA0144 gene product	BC003170	1.19
Cd642 effector protein 2	W81186	1.19
Consensus includes gb:AK023058.1 /DEF=Homo sapiens cDNA FLJ12996 fis, clone NTZRP3000235 /FEA=mRNA /DB_XREF=gi:10434801 /UG=Hs.287546 Homo sapiens	AX023058	1.19
hypothetical protein FLJ14345	NM_024733	1.19
gb:U50073.1 /DEF=Human T cell receptor beta chain precursor (Vbeta 12.3) mRNA, complete cds. /FEA=mRNA /GEN=Vbeta 12.3 /PROD=T cell receptor beta chain precursor	U50073	1.19
cadherin 2, type 1, N-cadherin (neuronal)	M34064	1.19
Consensus includes gb:AW517464 /FEA=EST /DB_XREF=gi:7155546 /DB_XREF=est:cp94g06.x1 /CLONE=IMAGE-2748058 /UG=Hs.119558 ribosomal protein L3	AW517464	1.19
hypothetical protein FLJ22222	NM_024848	1.19
casein kinase 1, epsilon	NM_001894	1.19
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10	AF060927	1.19
dehydrocratale dehydrogenase	M94065	1.19
palladin	AU157932	1.19
hypothetical protein FLJ13373	NM_025008	1.18
ELK3, ETS-domain protein (SRF accessory protein 2)	NM_005230	1.19
interleukin 18 receptor 1	NM_003855	1.19
bone morphogenetic protein 7 (osteogenic protein 1)	BC004248	1.19
lens intrinsic membrane protein 2 (19KD)	NM_030657	1.19
downregulated in ovarian cancer 1	NM_014890	1.19
squamous cell carcinoma antigen recognised by T cells 3	D63879	1.19
interleukin 2 receptor, alpha	K03122	1.19
KIAA0387 protein	AL138349	1.19
fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)	M63889	1.19
alkaline phosphatase, placental-like 2	J04946	1.19
microphthalmia-associated transcription factor	NM_000248	1.19
dishevelled, dish homolog 2 (Drosophila)	NM_004422	1.19
mucin 3A, intestinal	AF113816	1.19
PRO1748 protein	AF118073	1.19
hypothetical protein FLJ10636	NM_018279	1.19
deleted in lung and esophageal cancer 1	NM_007337	1.19
protocadherin alpha 2	BC003126	1.19
neuronal PAS domain protein 2	NM_002518	1.19
orosomucoid 1	NM_006607	1.19
hypothetical protein FLJ00001	AX024432	1.19
Consensus includes gb:AU146050 /FEA=EST /DB_XREF=gi:11007571 /DB_XREF=Hs.287459 Homo sapiens cDNA FLJ11111	AU146050	1.19
Tax1 (human T-cell leukemia virus type I) binding protein 1	U25801	1.19
KIAA0125 gene product	NM_014792	1.19
proopiomelanocortin (adrenocorticotropin/ beta-lipotropin/ alpha-melanocyte stimulating hormone/ beta-endorphin)	NM_000939	1.19
immune dysregulation, polyendocrinopathy, enteropathy, X-linked	NM_014009	1.19
beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)	NM_018644	1.19
Prader-Willi/Angelman syndrome-5	AU118874	1.19

protein kinase NYD-SP15
 KIA0583 gene product
 PRO1578 protein
 heparan sulfate (glucosamine) 3-O-sulfotransferase 1
 hypothetical protein FLJ23053
 CGI-101 protein
 Consensus includes gb:AF131844.1 /DEF=Homo sapiens clone 25015 mRNA /DB_XREF=gi:4406685 /UG=Hs.201628 Homo sapiens clone 25015 AF131844
 Consensus includes gb:AK024457.1 /DEF=Homo sapiens mRNA for FLJ00049 protein, partial cds. /FEA=mRNA /GEN=FLJ00049 /PROD=FLJ00049 protein /DB_XREF=gi:AK024457
 mitogen-activated protein kinase kinase 3
 leucine-rich repeat protein, neuronal 3
 ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)
 frequenin homolog (Drosophila)
 Kruppel-like factor 12
 transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)
 Notch homolog 2 (Drosophila)
 pregnancy specific beta-1-glycoprotein 1
 H4 histone family, member J
 retinol binding protein 4, plasma
 jerky homolog (mouse)
 glutamate receptor, metabotropic 7
 phospholamban
 sulfotransferase family, cytosolic, 2A, dehydroepiandrosterone (DHEA)-preferring, member 1
 sirutin silent mating type information regulation 2 homolog 2 (S. cerevisiae)
 forkhead box D1
 tropomyosin T3, skeletal, fast
 myeloid/lymphoid or mixed-lineage leukemia (littorax homolog, Drosophila)
 Consensus includes gb:AK028188.1 /DEF=Homo sapiens cDNA: FLJ22535 fls, clone HRC13115, highly similar to AF152336 Homo sapiens protocadherin gamma B7 (PCI AK028188)
 hypothetical protein PP1345
 KIAA1075 protein
 chondromodulin 1 precursor
 Consensus includes gb:AF070575.1 /DEF=Homo sapiens clone 24407 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387946 /UG=Hs.12432 Homo sapiens clone 24407 /AF070575
 T cell receptor alpha locus
 plakophilin 4
 nuclear receptor co-repressor 1
 paired basic amino acid cleaving system 4
 Rhesus blood group, D antigen
 tissue inhibitor of metalloproteinase 3 (Sorby fundus dystrophy, pseudoinflammatory)
 KIAA1609 protein
 hypothetical protein FLJ13285
 Consensus includes gb:A523613 /FEA=EST /DB_XREF=gi:4437748 /DB_XREF=est:ig95a07.x1 /CLONE=IMAGE:2118500 /UG=Hs.293495 ESTs, Weakly similar to ALU1
 pre-mRNA processing factor 18
 KIAA0863 protein
 myosin VI
 Consensus includes gb:l42230 /DEF=Homo sapiens core 2 beta-1,6-N-acetylglucosaminyltransferase pseudogene /FEA=CDS /DB_XREF=gi:1018163 /UG=Hs.247990 Ho L42230
 Consensus includes gb:AU148957 /FEA=EST /DB_XREF=gi:11070476 /DB_XREF=est:AU148957 /CLONE=NT2RM4001313 /UG=Hs.306638 Homo sapiens cDNA FLJ12: AU1543
 chromodomain helicase DNA binding protein 3
 small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated
 hypothetical protein FLJ10572
 hypothetical protein FLJ14033 similar to hypoxia inducible factor 3, alpha subunit
 microseminoprotein, beta-
 synrophin, beta 2 (dystrophin-associated protein A1, 59KD, basic component 2)
 PRY2 N-terminal domain-interacting receptor 1
 acid phosphatase, prostate
 small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (fractalkine, neurotactin)
 RAW binding protein 1
 integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)
 hypothetical protein FLJ20731
 uronyl 2-sulfotransferase
 ubiquitin-conjugating enzyme E2A (RAD6 homolog)
 hypothetical protein DKFZp434G2228
 DKFZP434K081 protein
 hypothetical protein FLJ20574
 integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)

tryptase beta 1
 Consensus includes gb:AK024909.1 /DEF=Homo sapiens cDNA: FLJ121256 fta, done COL01402. /FEA=mRNA /DB_XREF=gi:10437324 /UG=Hs.306724 Homo sapiens cC
 AK024909
 hypothetical protein FLJ12178
 peptide YY
 retinoic acid receptor, alpha
 CD3E antigen, epsilon polypeptide (TTT3 complex)
 Consensus includes gb:BE868287 /FEA=EST /DB_XREF=gi:10340242 /DB_XREF=est:60150987 /F1 /CLONE=IMAGE:3911808 /UG=Hs.249219 Homo sapiens chromoso
 interleukin 18 receptor accessory protein
 hypothetical protein FLJ12120
 cyclin, basic protein of sperm head cytoskeleton 1
 zinc-finger protein DZP1
 neuropeptide FF-amide peptide precursor
 adrenergic, beta, receptor kinase 2
 Consensus includes gb:AL353949.1 /DEF=Homo sapiens mRNA: cDNA DKFp761P1114 (from done DKFp761P1114). /FEA=mRNA /DB_XREF=gi:7669990 /UG=Hs.30
 telomeric repeat binding factor (NIMA-interacting) 1
 parvin, alpha
 small proline-rich protein 2C
 proboscoid gamma subfamily A, 3
 much 4, tracheobronchial
 nucleoporin 50K0
 serine/threonine kinase 38 like
 hypothetical protein DKFp566H0824
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 9
 zinc finger protein 237
 growth differentiation factor 9
 hypothetical protein
 KIAA0244 protein
 putative dipeptidase
 protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)
 sphingomyelin phosphodiesterase 2, neutral membrane (neutral sphingomyelinase)
 STAT induced STAT inhibitor-2
 hypothetical protein FLJ22662
 adrenomedullin receptor
 mahitin 2
 KIAA0459 protein
 PRO1900 protein
 bombesin-like receptor 3
 synovial sarcoma, X breakpoint 2
 Burkitt lymphoma receptor 1, GTP binding protein
 synaptoblagmin 1
 collagen, type VI, alpha 1
 reticulon 1
 hypothetical protein FLJ14529
 sh3-associated polypeptide, 30K0
 nuclear RNA export factor 2
 nuclear transcription factor Y, beta
 Consensus includes gb:AW574818 /FEA=EST /DB_XREF=gi:7246357 /DB_XREF=est:UI-HF-BK0-abi-e-06-0-UI.s1 /CLONE=IMAGE:3056362 /UG=Hs.32374 ESTs, Week
 venth 1
 phospholipase D2
 DKFZP586D0919 protein
 KIAA0871 protein
 armadillo repeat gene deletions in velocardiofacial syndrome
 Consensus includes gb:BF509345 /FEA=EST /DB_XREF=gi:11592709 /DB_XREF=est:UI-H-B14-aox-c-11-0-UI.s1 /CLONE=IMAGE:3086229 /UG=Hs.6996 ESTs
 glutathione S-transferase M4
 signal recognition particle receptor (docking protein)
 hypothetical protein FLJ12586
 coronin, actin binding protein, 2A
 phenylalanine-tRNA synthetase
 Lem3 protein
 PTK2 protein tyrosine kinase 2
 calcium channel, voltage-dependent, beta 2 subunit
 oxidised low density lipoprotein (lectin-like) receptor 1
 Consensus includes gb:AU144791 /FEA=EST /DB_XREF=gi:11006312 /DB_XREF=est:AU144791 /CLONE=HEMBA1002971 /UG=Hs.274237 Homo sapiens cDNA FLJ10
 AU144791

smoothened homolog (Drosophila) NM_005631 1.19
 small inducible cytokine A4 NM_002984 1.19
 insulin-like growth factor 2 (somatomedin A) NM_000612 1.19
 Consensus includes gb:AF052172.1 /DEF=Homo sapiens clone 24617.1 /AF052172 1.19
 protein (epiphy-prolyl isomerase) NIMA-interacting, 4 (osculin) NM_008223 1.19
 Consensus includes gb:A870151 /FEA=EST /DB_XREF=gi:5766983 /UG=Hs.77631 glycine cleavage system protei A870151 1.19
 hypothetical protein FLJ12606 NM_024604 1.19
 Consensus includes gb:AW138468 /FEA=EST /DB_XREF=gi:6142786 /DB_XREF=est:UH-B1-adtg-401-Q-U1.s1 /CLONE=IMAGE:2716920 /UG=Hs.299882 ESTs, Highly 1.19
 Consensus includes gb:AF090887.1 /DEF=Homo sapiens clone HQ0085 /FEA=mRNA /DB_XREF=gi:6690154 /UG=Hs.306562 Homo sapiens clone HQ0085 1.19
 spermatogenesis associated 2 NM_006038 1.19
 gamma-aminobutyric acid (GABA) B receptor, 1 AL031883 1.19
 Purkinje cell protein 4 NM_006198 1.19
 transforming growth factor, beta 2 M18154 1.19
 hypothetical protein FLJ20014 NM_017622 1.19
 glycoprotein hormones, alpha polypeptide NM_000735 1.19
 elastin (supraaortic stenosis, Williams-Beuren syndrome) M24782 1.19
 Consensus includes gb:AL121964 /DEF=Human DNA sequence from clone RP1-154G14 on chromosome 6q15-16.3. Contains the 3' end of the MAP3K7 gene for mitogen- 1.19
 cullin 4B AF212895 1.19
 mitogen-activated protein kinase 4 NM_002747 1.19
 Down syndrome cell adhesion molecule AF023450 1.19
 SRY (sex determining region Y)-box 20 NM_006942 1.19
 ribosomal protein S6 kinase, 52kD, polypeptide 1 NM_012424 1.19
 brain-derived neurotrophic factor NM_001709 1.19
 gamma-aminobutyric acid (GABA) A receptor, beta 2 NM_021911 1.19
 protein homologous to salivary proline-rich protein P-8 NM_012390 1.19
 G protein-coupled receptor 39 NM_001508 1.19
 Consensus includes gb:A1889513 /FEA=EST /DB_XREF=gi:5594677 /DB_XREF=est:wm05606.x1 /CLONE=IMAGE:2444578 /UG=Hs.168694 Homo sapiens clone 23763 u A1889513 1.19
 microtubule-associated protein, RPIEB family, member 3 BG222594 1.19
 globotriacylglyceramide/CO77 synthase; Gb3/CO77 synthase; alpha1.4-galactosyltransferase; 4-N-acetylglucosaminyltransferase NM_017436 1.19
 NEU1 protein NM_016215 1.19
 hypothetical protein BF787555 1.19
 non-canonical ubiquitin conjugating enzyme 1 AW500009 1.19
 SEC7 homolog NM_012455 1.19
 heat shock 27kD protein family, member 7 (cardiovascular) NM_014424 1.19
 Consensus includes gb:AK024879.1 /DEF=Homo sapiens cDNA: FLJ12226 fls, done COL00721. /FEA=mRNA /DB_XREF=gi:10437291 /UG=Hs.306715 Homo sapiens cDNA AK024879 1.19
 putative protein-tyrosine kinase NM_015978 1.19
 Consensus includes gb:AF056433.1 /DEF=Homo sapiens clone FBD3 Cn-du-chat critical region mRNA. /FEA=mRNA /DB_XREF=gi:3044151 /UG=Hs.8078 Homo sapiens 1.19
 hypothetical protein MGC5350 NM_030920 1.19
 zinc finger protein 26 (KOX 20) NM_019591 1.19
 multiple endocrine neoplasia 1 NM_000244 1.19
 protein-L-isoaspartate (D-aspartate) O-methyltransferase NM_005389 1.19
 M11167 1.19
 Consensus includes gb:AK022309.1 /DEF=Homo sapiens cDNA FLJ12247 fls, clone MAMMA1001397. /FEA=mRNA /DB_XREF=gi:10433678 /UG=Hs.287506 Homo sapiens 1.19
 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 2 W72516 1.19
 Consensus includes gb:AA102667 /FEA=EST /DB_XREF=gi:1647984 /DB_XREF=est:zn73e05.s1 /CLONE=IMAGE:563840 /UG=Hs.284673 ESTs, Weakly similar to ALU8 1.19
 Consensus includes gb:AA102667 /FEA=EST /DB_XREF=gi:1647984 /DB_XREF=est:zn73e05.s1 /CLONE=IMAGE:563840 /UG=Hs.284673 ESTs, Weakly similar to ALU8 1.19
 glypican 4 NM_001448 1.19
 hypothetical protein NM_019103 1.19
 adaptor-related protein complex 3, sigma 2 subunit BC002785 1.19
 Consensus includes gb:BE046461 /FEA=EST /DB_XREF=gi:8363514 /DB_XREF=est:hm47g05.x2 /CLONE=IMAGE:3026840 /UG=Hs.159412 Homo sapiens clone 24628 r BE046461 1.19
 hypothetical protein FLJ20075 NM_017655 1.19
 fasciculation and elongation protein zeta 1 (zyglin I) NM_022549 1.19
 cytochrome P450, subfamily I1C (mephenytoin 4-hydroxylase), polypeptide 8 NM_030878 1.19
 G protein-coupled receptor 9 NM_001504 1.19
 hypothetical protein FLJ20005 NM_017617 1.19
 hypothetical protein FLJ10865 NM_018173 1.19
 RAB11B, member RAS oncogene family X78780 1.19
 mitogen-activated protein kinase kinase 13 NM_004721 1.19
 dynein, axonemal, intermediate polypeptide, 1 NM_012144 1.19
 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) M69176 1.19
 tropoin T2, cardiac X79857 1.19
 tumor suppressing subtransferable candidate 3 BC005034 1.19
 calmeglin NM_004362 1.19

potassium inwardly-rectifying channel, subfamily J, member 3
hypothetical protein FLJ20958
paired box gene 4
Consensus includes gb:AL520875 /FEA=EST /DB_XREF=gi:12784168 /DB_XREF=est:AL520875 /CLONE=CS0DB002YF15 (3 prime) /UG=Hs.77783 membrane-associate AL520875
Consensus includes gb:AF141801.1 /DEF=Homo sapiens GKAPSAPAP interacting protein (SHANK) mRNA, partial cds. /FEA=mRNA /GEN=SHANK /PROD=GKAPSAPAF AF141801
Consensus includes gb:AF141801.1 /DEF=Homo sapiens GKAPSAPAP interacting protein (SHANK) mRNA, partial cds. /FEA=mRNA /GEN=SHANK /PROD=GKAPSAPAF AF141801
hypothetical protein FLJ11186
platelet factor 4 variant 1
senna domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3F
ribosome binding protein 1 homolog 180kD (dog)
sperm protein associated with the nucleus, X chromosome, family member A1
hypothetical protein FLJ11183
alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae)
claudin 14
alcohol dehydrogenase 1A (class I), alpha polypeptide
cell division cycle 25B
hypothetical protein F23149_1
fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group included)
Consensus includes gb:AA417256 /FEA=EST /DB_XREF=gi:2077373 /DB_XREF=est:zu07e04.s1 /CLONE=IMAGE:731166 /UG=Hs.193769 ESTs
KJAA0349 protein
hypothetical protein FLJ10282
cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kD
guanine nucleotide binding protein-like 1
integral inner nuclear membrane protein
sperm associated antigen 6
prostaglandin E receptor 1 (subtype EP1), 42kD
Consensus includes gb:AV700288 /FEA=EST /DB_XREF=gi:10302269 /DB_XREF=est:AV700288 /CLONE=GKCBVG05 /UG=Hs.285173 ESTs, Moderately similar to CD4 AV700288
nucleoporin 98kD
Consensus includes gb:AF070810.1 /DEF=Homo sapiens clone 24505 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387982 /UG=Hs.100543 Homo sapiens clone 24505 AF070810
hypothetical protein FLJ10634
hypothetical protein MGC10783
suppressor of S. cerevisiae gcr2
nuclear receptor subfamily 4, group A, member 1
neuronal pentraxin II
Consensus includes gb:AU154785 /FEA=EST /DB_XREF=gi:1016308 /DB_XREF=est:AU154785 /CLONE=NT2RP4002888 /UG=Hs.31532 Homo sapiens mRNA; cDNA (AU154785
RAD52 homolog (S. cerevisiae)
hypothetical protein DKFZp566M1120
DKFZp566M0923 protein
profilin 2
serine protease inhibitor, Kazal type 2 (acrosin-lypsin inhibitor)
centrin, EF-hand protein, 3 (CDC31 homolog, yeast)
gamma-aminobutyric acid (GABA) A receptor, alpha 4
Consensus includes gb:BC000975.2 /DEF=Homo sapiens, clone IMAGE:3448308, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3448308) /DB_XI BC000975
CD84 antigen (leukocyte antigen)
hypothetical protein FLJ22757
hypothetical protein FLJ11729
DKFZP566A0946 protein
actinin, alpha 1
calpain 2, (mu) large subunit
Fc fragment of IgA, receptor for
SEC24 related gene family, member B (S. cerevisiae)
protein tyrosine phosphatase, non-receptor type 9
solute carrier family 12 (sodium/chloride transporters), member 3
alkylation repair, alkB homolog
necroline receptor 2 (cardiac)
small inducible cytokine subfamily A (Cys-Cys), member 13
ataxia telangiectasia mutated (includes complementation groups A, C and D)
clone FLB5214
intercellular adhesion molecule 5, telencephalin
solute carrier family 7, cationic amino acid transporter, y+ system) member 10
Consensus includes gb:AK027217.1 /DEF=Homo sapiens cDNA: FLJ23564 fls, clone LNG10773, /FEA=mRNA /DB_XREF=gi:10440296 /UG=Hs.306913 Homo sapiens cDNA: AK027217
transglutaminase 5
hypothetical protein FLJ20507
nucleolar protein family A, member 2 (HACA small nucleolar RNPs)

nuclear receptor coactivator 3	NM_006534	1.18
Consensus includes gb:AK025177.1 /DEF=Homo sapiens cDNA: FLJ21524 fts, clone COL05921 /FEA=mRNA /DB_XREF=gi:10437640 /UG=Hs.306778 Homo sapiens cDNA: AK025177	NM_000116	1.18
tariazin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome)	NM_004375	1.18
COX11 homolog, cytochrome c oxidase assembly protein (yeast)	NM_000900	1.18
matrix Gla protein	NM_024732	1.18
hypothetical protein FLJ14351	NM_004670	1.18
3'-phosphoadenosine 5'-phosphosulfate synthase 2	NM_005832	1.18
potassium large conductance calcium-activated channel, subfamily M, beta member 2	NM_005399	1.18
protein kinase, AMP-activated, beta 2 non-catalytic subunit	AV700403	1.18
Consensus includes gb:AV700403 /FEA=EST /DB_XREF=gi:10302374 /DB_XREF=est:AV700403 /CLONE=GK6ADA01 /UG=Hs.293673 ESTs	NM_003201	1.18
transcription factor A, mitochondrial	AJ246641	1.18
hypothetical protein FLJ14154	NM_025163	1.18
hypothetical protein FLJ12768	BF509089	1.18
hypothetical protein MGC20727	AL109874	1.18
Consensus includes gb:AL109874.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 46866 /FEA=mRNA /DB_XREF=gi:5688832 /UG=Hs.221811	NM_020250	1.18
MOST2 protein	NM_024706	1.18
hypothetical protein FLJ13479	NM_022912	1.18
hypothetical protein FLJ13110	NM_003645	1.18
fatty-acid-Coenzyme A ligase, very long-chain 1	NM_002935	1.18
ribonuclease, RNase A family, 3 (eosinophil cationic protein)	AF265167	1.18
ATP-binding cassette, sub-family A (ABC1), member 1	AF239764	1.18
EGF-like module-containing mucin-like receptor EMR3	NM_000810	1.18
gamma-aminobutyric acid (GABA) A receptor, alpha 5	NM_015880	1.18
RIG-like 14-1	NM_021235	1.18
epidermal growth factor receptor substrate EPS15R	AI358867	1.18
apolipoprotein E	AI837543	1.18
DC12 protein	NM_001881	1.18
epidermal growth factor receptor pathway substrate 15	AC004755	1.18
one cut domain, family member 3	AI863043	1.18
Consensus includes gb:AI863043 /FEA=EST /DB_XREF=gi:5810262 /DB_XREF=est:hw320b11.x1 /CLONE=IMAGE:2359549 /UG=Hs.246885 hypothetical protein FLJ2078	AL049242	1.18
Consensus includes gb:AL049242.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564B083 (from clone DKFZp564B083) /FEA=mRNA /DB_XREF=gi:4499977 /UG=Hs.2347	AV651241	1.18
Consensus includes gb:AV651241 /FEA=EST /DB_XREF=gi:9872255 /DB_XREF=est:AV651241 /CLONE=GLCCNG03 /UG=Hs.282469 ESTs	AA046411	1.18
amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AF188773	1.18
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 3	NM_004636	1.18
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B	U86584	1.18
crystallin, alpha A	AF109294	1.18
methylthadenosine phosphorylase	AK022102	1.18
Consensus includes gb:AK022102.1 /DEF=Homo sapiens cDNA FLJ12040 fts, clone HEMMB1001944 /FEA=mRNA /DB_XREF=gi:10433423 /UG=Hs.296687 Homo sapiens cDNA: AK022102	NM_008071	1.18
polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jelly, sea urchin homolog) like	NM_005380	1.18
v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian)	AJ796813	1.18
schillin	NM_005048	1.18
parathyroid hormone receptor 2	AL046884	1.18
retinoblastoma (X-linked, juvenile) 1	L00692	1.18
carcinoembryonic antigen-related cell adhesion molecule 3	AF007141	1.18
5-hydroxytryptamine (serotonin) receptor 6	AW268840	1.18
eukaryotic translation initiation factor 4E	NM_005756	1.18
G protein-coupled receptor 64	U94506	1.18
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)	AK023069	1.18
hypothetical protein	AL574660	1.18
ATP-binding cassette, sub-family D (ALD), member 4	NM_022820	1.18
cytochrome P450 polypeptide 43	AW026846	1.18
C1q-related factor	NM_002939	1.18
ribonuclease/angiogenin inhibitor	AI023317	1.18
KIAA0130 gene product	NM_024582	1.18
hypothetical protein FLJ23056	NM_023922	1.18
taste receptor, type 2, member 14	NM_006355	1.18
ring finger protein 15	AI341811	1.18
gene amplified in squamous cell carcinoma 1	NM_001386	1.18
dihydropyrimidinase-like 2	X88405	1.18
myelin associated glycoprotein	U57001	1.18
ephrin-B3	NM_006890	1.18
carcinoembryonic antigen-related cell adhesion molecule 7	NM_002585	1.18
pre-B-cell leukemia transcription factor 1		

Consensus includes gb:AK023816.1 /DEF=Homo sapiens cDNA FLJ13754 fls, clone PLACE3000362. /FEA=mRNA /DB_XREF=gj:10435965 /UG=Hs.136233 Homo sapiens inhibin, alpha MI3881 1.18

NIMA (never in mitosis gene a)-related kinase 1 AF155113 1.18

peroxisome proliferative activated receptor, alpha NM_005036 1.18

MAD, mothers against decapentaplegic homolog 6 (Drosophila) AF035528 1.18

hypothetical protein PP1628 NM_025201 1.18

hypothetical protein DKFZP434C135 NM_030938 1.18

ubiquitously transcribed tetrapeptide repeat gene, Y chromosome AF000895 1.18

lucosyltransferase 6 (alpha 1.3) lucosyltransferase UZ7335 1.18

cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6 AF182275 1.18

AD037 protein AI890191 1.18

FXYD domain-containing ion transport regulator 7 NM_022006 1.18

KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3 NM_016857 1.18

MHC class I region ORF NM_006674 1.18

TGFB inducible early growth response 2 AA149594 1.18

hypothetical protein FLJ23259 NM_024727 1.18

J domain containing protein 1 NM_021800 1.18

sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F NM_004263 1.18

sperm protein associated with the nucleus, X chromosome, family member A1 NM_005162 1.18

Consensus includes gb:BF344237 /FEA=EST /DB_XREF=gj:1291450 /DB_XREF=est:60201749 /f1 /CLONE=IMAGE:4152983 /UG=Hs.124776 Homo sapiens mRNA, cc BF344237 BE964082 1.18

complement component 1, q subcomponent binding protein NM_005312 1.18

guanine nucleotide-releasing factor 2 (specific for crk proto-oncogene) A1816291 1.18

hypothetical protein BF440025 1.18

nephroblastoma overexpressed gene gb:AF165520.1 /DEF=Homo sapiens phorbol 1 protein (PBI) mRNA, complete cds. /FEA=mRNA /GEN=PBI /PROD=phorbol 1 protein /DB_XREF=gj:9294746 /UG=Hs.85 AF165520 1.18

astrotactin AB006627 1.18

ATPase, Class I, type 8B, member 2 AL137537 1.18

hypothetical protein MGC4659 NM_025268 1.18

collagen, type V, alpha 2 AL575735 1.18

interleukin 4 NM_000589 1.18

helicase-moi AK001827 1.18

histamine receptor H1 D28481 1.18

Consensus includes gb:BE965029 /FEA=EST /DB_XREF=gj:11768976 /DB_XREF=est:601658812R1 /CLONE=IMAGE:3886131 /UG=Hs.198793 Homo sapiens cDNA, FL BE965029 1.18

KIAA1654 protein AB051441 1.18

Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse) NM_002392 1.18

fasciculation and elongation protein zeta 2 (zyglin II) L17328 1.18

CEGP1 protein A1424243 1.18

interferon induced transmembrane protein 1 (9-27) NM_003641 1.18

B-cell CLL/lymphoma 11B (zinc finger protein) NM_022898 1.18

MUF1 protein AK024051 1.18

v-rel reticuloendotheliosis viral oncogene homolog A, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3, p65 (avian) NM_021975 1.18

fibroblast growth factor 18 BC008245 1.18

leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2 AF004231 1.18

proteolipid protein 1 (Pelizaeus-Merzbacher disease, spastic paraplegia 2, uncomplicated) BC002665 1.18

protein O-acyltransferase 1 AL045513 1.18

advanced glycosylation end product-specific receptor AB036432 1.18

A kinase (PKA) anchor protein (yoliao) 9 NM_005751 1.18

NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) pseudogene 5 AL050308 1.18

tripartite motif-containing 32 BC003154 1.18

ectodermal dysplasia 1, anhidrotic EC001194 1.18

nucleotide-sugar transporter similar to C. elegans sqv-7 AF081194 1.18

2',3'-cyclic nucleotide 3' phosphodiesterase AJ005866 1.18

gb:136759.1 /DEF=Human pre-T cell receptor alpha-type chain precursor, mRNA, complete cds. /FEA=mRNA /PROD=pre-T cell receptor alpha-type chain precursor /DB_XREF=gj:93759 1.18

replication initiation region protein (60kD) NM_014374 1.18

hairly homolog (Drosophila) BE973687 1.18

phosphatidate-3-kinase, class 3 NM_022647 1.18

rab3 interacting protein AF263310 1.18

erythrocyte transmembrane protein NM_016158 1.18

KIAA1098 protein A1359472 1.18

tubulin, beta, 2 AA515698 1.18

myelin associated glycoprotein X98405 1.18

v-erb-a erythroblastic leukemia viral oncogene homolog 4 (avian) NM_005235 1.18

drebrin 1
 CDC-like kinase 1
 bone morphogenetic protein 4
 ribosomal protein L34 pseudogene 2
 Consensus includes gb:BE856707 /FEA=EST /DB_XREF=gi:10370006 /DB_XREF=est:7166102.x1 /CLONE=IMAGE:3299643 /UG=Hs.12520 Homo sapiens clone 23568.2 BE856707
 KIAA0334 protein
 homeo box B13
 CD58 antigen, (lymphocyte function-associated antigen 3)
 transmembrane 4 superfamily member 2
 collagen, type I, alpha 1
 Consensus includes gb:A1978930 /FEA=EST /DB_XREF=gi:5804020 /DB_XREF=est:wr68h11.x1 /CLONE=IMAGE:2492901 /UG=Hs.125258 Homo sapiens cDNA FLJ1371 A1978930
 Hypothetical protein FLJ20452
 Huntingtin interacting protein B
 follicle stimulating hormone, beta polypeptide
 carboxylesterase 2 (intestine, liver)
 PR00149 protein
 hypothetical protein FLJ12735
 protein phosphatase 1, regulatory (inhibitor) subunit 9A
 neuroepithelial cell transforming gene 1
 BUB1 budding uninhibited by benzimidazoles 1 homolog beta (yeast)
 roundsabout, axon guidance receptor, homolog 1 (Drosophila)
 M-phase phosphoprotein 9
 Cdc42 effector protein 2
 troponin C2, fast
 hypothetical protein FLJ10040
 KIAA0370 protein
 pater pan homolog (Drosophila)
 KIAA0470 gene product
 sonic hedgehog homolog (Drosophila)
 ribosomal protein S28
 Consensus includes gb:BC001878.1 /DEF=Homo sapiens, clone IMAGE:3534181, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3534181) /DB_XREF=est:yd3b02.s1 /CLONE=IMAGE:24533 /UG=Hs.155959 Homo sapiens clone 24533 mRNA R38712
 hypothetical protein FLJ13639
 fibrinogen, A alpha polypeptide
 uncharacterized hypothalamus protein HT010
 matrix metalloproteinase 24 (membrane-inserted)
 hypothetical protein FLJ22476
 hypothetical protein FLJ20208
 ammonites protein
 Consensus includes gb:R38712 /FEA=EST /DB_XREF=gi:796168 /DB_XREF=est:yd3b02.s1 /CLONE=IMAGE:24533 /UG=Hs.155959 Homo sapiens clone 24533 mRNA R38712
 6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 2
 C1q-related factor
 CGI-96 protein
 Consensus includes gb:AB051440.1 /DEF=Homo sapiens mRNA for KIAA1653 protein, partial cds. /FEA=mRNA /GEN=KIAA1653 /PROD=KIAA1653 protein /DB_XREF=gi:AB051440
 plectestin
 T-box 3 (lunar mammary syndrome)
 v-Jun sarcoma virus 17 oncogene homolog (avian)
 apolipoprotein D
 hypothetical protein FLJ20259
 hypothetical protein FLJ20363
 bridging integrator 1
 serine/threonine kinase FKSG61
 sodium channel, voltage-gated, type II, alpha 2 polypeptide
 Consensus includes gb:U92992.1 /DEF=Homo sapiens clone DT1PIA11 mRNA, CAG repeat region. /FEA=mRNA /DB_XREF=gi:2781410 /UG=Hs.168303 Homo sapiens AF103574
 immunoglobulin kappa constant
 RAB4, member RAS oncogene family
 NEL-like 2 (chicken)
 synovial sarcoma, X breakpoint 2
 much 2, intestinal/tracheal
 solute carrier family 22 (organic anion transporter), member 8
 KIAA0841 protein

Consensus includes gb:AL442084.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547G044 (from clone DKFZp547G044). /FEA=mRNA /DB_XREF=gi:10241763 /UG=Hs.306 AL442084
 G protein-coupled receptor 52 NM_005684 1.18
 KIA00750 gene product NM_014632 1.18
 ATP-binding cassette, sub-family G (WHITE), member 2 AF089951 1.18
 ras responsive element binding protein 1 AW118862 1.18
 Consensus includes gb:AK025086.1 /DEF=Homo sapiens cDNA: FLJ21433 fs, clone COL04232. /FEA=mRNA /DB_XREF=gi:10437530 /UG=Hs.306761 Homo sapiens cDNA AK025086
 homeo box C13 NM_017410 1.18
 HYA22 protein BF031714 1.18
 calcium channel, voltage-dependent, beta 2 subunit NM_000724 1.18
 adhesion glycoprotein NM_005666 1.18
 H2B histone family, member D NM_003520 1.18
 adenomatous polyposis coli NM_000038 1.18
 a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 12 NM_030955 1.18
 adrenergic, alpha-2A-, receptor AF284095 1.18
 calcium binding protein 2 NM_016366 1.18
 amphiregulin (schwannoma-derived growth factor) NM_001657 1.18
 annexin A8 NM_001630 1.18
 H1 histone family, member 4 AL353759 1.18
 ribonucleotide reductase M1 polypeptide NM_001033 1.18
 Consensus includes gb:AI590869 /FEA=EST /DB_XREF=gi:4599517 /DB_XREF=est:tw88g03.x1 /CLONE=IMAGE:2265804 /UG=Hs.172035 hypothetical protein similar to AI590869
 H3 histone, family 3A A1955855 1.18
 indoleamine-pyrole 2,3 dioxygenase M34455 1.18
 Consensus includes gb:AL137713.1 /DEF=Homo sapiens mRNA; cDNA DKFZp4340523 (from clone DKFZp4340523). /FEA=mRNA /DB_XREF=gi:6808127 /UG=Hs.3064 AL137713
 KIA00887 protein AB020694 1.18
 period homolog 1 (Drosophila) NM_002616 1.18
 mab-21-like 2 (C. elegans) AF262032 1.18
 heat shock 70kD protein 8 (HSP70B) NM_002155 1.18
 hypothetical protein FLJ11040 BF688108 1.18
 forkhead box Q3A AF041336 1.18
 Consensus includes gb:AL049213.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564B076 (from clone DKFZp564B076). /FEA=mRNA /DB_XREF=gi:4500088 /UG=Hs.2110: AL049213
 Consensus includes gb:BG149837 /FEA=EST /DB_XREF=gi:12661867 /DB_XREF=est:nae01b12.x1 /CLONE=IMAGE:3434082 /UG=Hs.33540 EST's, Weakly similar to dL: BG149837
 Consensus includes gb:Z25428.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete cds /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405734 /FL=gb:Z25428
 hepatocellular carcinoma antigen gene 520 NM_022087 1.18
 hypothetical protein CDA08 NM_030790 1.18
 MyoD family inhibitor NM_005586 1.18
 Consensus includes gb:LA8728 /DEF=Homo sapiens T cell receptor beta (TCRBV10S1) gene, complete cds /FEA=mRNA_1 /DB_XREF=gi:1054550 /UG=Hs.37163 Homo LA8728
 Testis-specific XK-related protein on Y NM_004877 1.18
 gb:Z25428.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete cds /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405734 /FL=gb:Z25428
 KIA00650 protein NM_0014550 1.18
 myosin IXA NM_006901 1.18
 KIA00800 gene product NM_014703 1.18
 hypothetical protein FLJ10210 NM_018027 1.18
 small inducible cytokine B subfamily (Cys-X-Cys motif), member 13 (B-cell chemoattractant) NM_006419 1.17
 small proline-rich protein 18 (cornalin) NM_003125 1.17
 surfactant, pulmonary-associated protein C N56912 1.17
 alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae) BC002521 1.17
 fetuin B AB017551 1.17
 Consensus includes gb:AL137428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N1323 (from clone DKFZp761N1323). /FEA=mRNA /DB_XREF=gi:6807885 /UG=Hs.30 AL137428
 Consensus includes gb:AL049988.1 /DEF=Homo sapiens mRNA; cDNA DKFZp584F212 (from clone DKFZp584F212). /FEA=mRNA /DB_XREF=gi:4884239 /UG=Hs.3063 AL049988
 interleukin 11 receptor, alpha NM_003277 1.17
 claudin 5 (transmembrane protein deleted in velocardiofacial syndrome) NM_004512 1.17
 Consensus includes gb:AL117652.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1324 (from clone DKFZp586B1324). /FEA=mRNA /DB_XREF=gi:5912244 /UG=Hs.271 AL117652
 pan-hematopoietic expression NM_005705 1.17
 angiotensin I converting enzyme (peptidyl-dipeptidase A) 2 NM_021804 1.17
 KIA00321 protein AB002319 1.17
 chromosome 8 open reading frame 5 AW296296 1.17
 PR domain containing 13 NM_021620 1.17
 basic leucine zipper nuclear factor 1 (JEM-1) AF286161 1.17
 Huntingtin interacting protein H AF161412 1.17
 KIA00323 protein A1075450 1.17
 ATP-binding cassette, sub-family D (ALD), member 2 NM_005164 1.17
 KIA00471 gene product AB019480 1.17
 T-cell leukemia/lymphoma 6 NM_020552 1.17
 hypothetical protein FLJ20909 NM_024693 1.17

hypothetical gene CG030 1.17 U50531
 inducible T-cell co-stimulator 1.17 AB023135
 protein kinase (CAMP-dependent, catalytic) inhibitor gamma 1.17 NM_007066
 CGI-121 protein 1.17 NM_016058
 Consensus includes gp:AL049992.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G242 (from clone DKFZp564G242) /FEA=mRNA /DB_XREF=gi:4884244 /UG=Hs.2258 /AL049992 1.17
 glypican 3 1.17 L47125
 hypothetical protein MGC3146 1.17 NM_031289
 myosin IXb 1.17 NM_004145
 hepatocyte nuclear factor 3, alpha 1.17 NM_004496
 matrix metalloproteinase 11 (stromelysin 3) 1.17 AA0401885
 keratin 12 (Meesmann corneal dystrophy) 1.17 NM_000223
 crystallin, gamma B 1.17 NM_005210
 dipeptidylpeptidase IV (CD26, adenosine deaminase complementing protein 2) 1.17 NM_001835
 X-ray repair complementing defective repair in Chinese hamster cells 4 1.17 BC052569
 pleckstrin homology, Sec7 and coiled/coiled domains 4 1.17 NM_013385
 Consensus includes gp:AA47740 /FEA=EST /DB_XREF=gi:2161410 /DB_XREF=est:aa1905.s1 /CLONE=IMAGE.813633 /UG=Hs.99139 Homo sapiens cDNA: FLJ2294f:AA447740 1.17
 phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 duncie homolog, Drosophila) 1.17 L20966
 fatty acid binding protein 6, ileal (gastrotopin) 1.17 U19869
 translocase of outer mitochondrial membrane 70 homolog A (yeast) 1.17 NM_014820
 complement component 3a receptor 1 1.17 U62027
 gb:NM_005405.1 /DEF=Homo sapiens RNA binding motif protein, Y chromosome, family 2, member B (RBMV2B) mRNA. /FEA=mRNA /GEN=RBMV2B /PROD=RNA bindi 1.17
 hypothetical protein FLJ23119 1.17 NM_024652
 14-3-3 protein 1.17 A1999677
 neural cell adhesion molecule 1 1.17 AA128505
 KIAA0130 gene product 1.17 AK022508
 Consensus includes gp:AW340588 /FEA=EST /DB_XREF=gi:6837214 /DB_XREF=est:hd5a05.x1 /CLONE=IMAGE.2906592 /UG=Hs.8736 Homo sapiens mRNA; cDNA 1.17
 heat shock protein (hsp110 family) 1.17 NM_014278
 hypothetical protein FLJ14393 1.17 BE501352
 cer-d4 (mouse) homolog 1.17 NM_012074
 spectre-type POZ protein 1.17 BF873888
 ras association (Raf/GDS/AF-6) domain containing protein JC265 1.17 AL136924
 methionine adenosyltransferase II, beta 1.17 NM_013283
 Consensus includes gp:AB84887 /FEA=EST /DB_XREF=gi:5590031 /DB_XREF=est:w85g03.x1 /CLONE=IMAGE.2431732 /UG=Hs.91379 ribosomal protein L26 1.17
 heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A) 1.17 A1884867
 KIAA0275 gene product 1.17 D13413
 neuropilin 2 1.17 NM_014767
 PROO159 protein 1.17 AF280546
 tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein) 1.17 NM_014118
 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B 1.17 AF028071
 putative leukocyte platelet-activating factor receptor 1.17 NM_004636
 Consensus includes gp:Y10255.1 /DEF=H.sapiens ACTH receptor mRNA 3UTR. /FEA=mRNA /DB_XREF=gi:2230860 /UG=Hs.274501 H.sapiens ACTH receptor mRNA 3' 1.17
 soluble carrier family 21 (organic anion transporter), member 3 1.17 NM_022571
 cytotoxic T-lymphocyte-associated protein 4 1.17 AF085224
 SH3-domain GRB2-like 3 1.17 NM_005214
 Consensus includes gp:AJ08433 /DEF=Homo sapiens partial IGKV gene for immunoglobulin kappa chain variable region, clone 38 /FEA=CDS /DB_XREF=gi:12655549 1.17
 sine oculis homeobox homolog 2 (Drosophila) 1.17 NM_016932
 gap junction protein, beta 1, 32kD (connexin 32, Charcot-Marie-Tooth neuropathy, X-linked) 1.17 NM_000166
 DKFZP564I052 protein 1.17 AL080063
 Consensus includes gp:AL137284.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D1516 (from clone DKFZp434D1516) /FEA=mRNA /DB_XREF=gi:8807739 /UG=Hs.27 1.17
 tumor protein 63 kDa with strong homology to p53 1.17 AL137284
 zinc-finger protein ZBRK1 1.17 AB042841
 histone acetyltransferase 1 family, member A3 1.17 NM_021632
 gap junction protein, alpha 5, 40kD (connexin 40) 1.17 NM_012330
 aldohyde dehydrogenase 1 family, member A3 1.17 NM_000693
 retinoid X receptor, gamma 1.17 NM_006917
 keratohelin alpha 1 (importin alpha 5) 1.17 AW051311
 protodactherin 22 1.17 AJ276803
 KIAA0852 gene product 1.17 AU153525
 cathepsin H 1.17 NM_004390
 kiulonidase, alpha-L- 1.17 M74715
 Consensus includes gp:AL520908 /FEA=EST /DB_XREF=gi:12784401 /DB_XREF=est:AL520908 /CLONE=CS0D9002YF01 (3 prime) /UG=Hs.155489 NS1-associated prc 1.17
 malonyl-CoA decarboxylase 1.17 NM_012213
 gap junction protein, alpha 5, 40kD (connexin 40) 1.17 NM_005266
 tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor) 1.17 AF021834

SET binding protein 1	NM_015559	1.17
hypothetical protein PRO3073	NM_018549	1.17
chromosome 22 open reading frame 4	AL096779	1.17
casease 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	U13700	1.17
platelet-activating factor acetylhydrolase 2 (40kD)	U89386	1.17
KUAA0420 gene product	AB007880	1.17
uncharacterized hypothalamus protein HSNINP1	NM_018478	1.17
X04603 B subunit bnc, bnc genes corresponding to nucleotides 248-2229 of X04603 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	X04603	1.17
eryl hydrocarbon receptor	NM_001621	1.17
hypothetical protein FLJ110971	NM_018287	1.17
target of myb1-like 1 (chicken)	gb:NM_016425.1 / DEF=Homo sapiens transmembrane protease, serine 4 / (TMPRSS4), mRNA, / FEA=mRNA / GEN=TMPRSS4 / PROD=transmembrane protease, serine 4 / NM_016425	1.17
Consensus includes gb:BE672818 / FEA=EST / DB_XREF=gi:10033359 / DB_XREF=est:7d24c01.x1 / CLONE=IMAGE:3248160 / JG=Hs.297681 serine (or cysteine) protease BE672818	NM_005486	1.17
Consensus includes gb:BE672818 / FEA=EST / DB_XREF=gi:10033359 / DB_XREF=est:7d24c01.x1 / CLONE=IMAGE:3248160 / JG=Hs.297681 serine (or cysteine) protease BE672818	NM_030820	1.17
collagen, type XXI, alpha 1	NM_003657	1.17
Consensus includes gb:BE672818 / FEA=EST / DB_XREF=gi:10033359 / DB_XREF=est:7d24c01.x1 / CLONE=IMAGE:3248160 / JG=Hs.297681 serine (or cysteine) protease BE672818	U05255	1.17
breast carcinoma amplified sequence 1	NM_018231	1.17
glycophorin B (includes Ss blood group)	NM_022349	1.17
hypothetical protein FLJ10815	NM_000060	1.17
membrane-spanning 4-domains, subfamily A, member 6A	NM_003995	1.17
Consensus includes gb:AL035250 / DEF=Human DNA sequence from clone RP4-614C15 on chromosome 20q13.2-13.33 Contains the EDN3 gene for endothelin 3, ESTs, AL035250	NM_014191	1.17
Consensus includes gb:AL035250 / DEF=Human DNA sequence from clone RP4-614C15 on chromosome 20q13.2-13.33 Contains the EDN3 gene for endothelin 3, ESTs, AL035250	NM_012465	1.17
Consensus includes gb:AL035250 / DEF=Human DNA sequence from clone RP4-614C15 on chromosome 20q13.2-13.33 Contains the EDN3 gene for endothelin 3, ESTs, AL035250	NM_006285	1.17
Consensus includes gb:AL117481.1 / DEF=Homo sapiens mRNA; cDNA DKFZp547F134 (from clone DKFZp547F134), / FEA=mRNA / DB_XREF=gi:12224853 / JG=Hs.147: AL117481	BF665566	1.17
CD58 antigen, (lymphocyte function-associated antigen 3)	BF665566	1.17
ras-related C3 botulinum toxin substrate 3 (rho family, small GTP binding protein Rac3)	NM_001779	1.17
bioidkase	NM_005052	1.17
natriuretic peptide receptor Bguanylate cyclase B (atrial natriuretic peptide receptor B)	NM_000060	1.17
sodium channel, voltage gated, type VIII, alpha polypeptide	NM_003995	1.17
gb:NM_012465.1 / DEF=Homo sapiens tctid-like 2 (TLL2), mRNA, / FEA=mRNA / GEN=TL2 / PROD=tctid-like 2 / DB_XREF=gi:6912723 / FL=gb:NM_012465.1	NM_014191	1.17
testis-specific kinase 1	NM_012465	1.17
Consensus includes gb:AK026856.1 / DEF=Homo sapiens cDNA: FLJ23203 fis, clone AKA02487, / FEA=mRNA / DB_XREF=gi:10439812 / JG=Hs.306889 Homo sapiens c AK026856	NM_001779	1.17
leucine rich repeat (in FLN) interacting protein 1	BF665566	1.17
Consensus includes gb:AL049443.1 / DEF=Homo sapiens mRNA; cDNA DKFZp568N2020 (from clone DKFZp568N2020), / FEA=mRNA / DB_XREF=gi:4500224 / JG=Hs.16 AL049443	NM_014256	1.17
Consensus includes gb:AL049443.1 / DEF=Homo sapiens mRNA; cDNA DKFZp568N2020 (from clone DKFZp568N2020), / FEA=mRNA / DB_XREF=gi:4500224 / JG=Hs.16 AL049443	U08024	1.17
UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 3	A815793	1.17
sulfotransferase family, cytosolic, 2A, dehydroepiandrosterone (DHEA)-preferring, member 1	NM_019857	1.17
protein kinase C substrate 80K-H	NM_001442	1.17
CTP synthase II	AB017915	1.17
fatty acid binding protein 4, adipocyte	AK022067	1.17
carbohydrate (chondroitin & keratan) sulfotransferase 3	A184136	1.17
Consensus includes gb:AK022067.1 / DEF=Homo sapiens cDNA FLJ12005 fis, clone HEMB81001565, / FEA=mRNA / DB_XREF=gi:10433387 / JG=Hs.287477 Homo sapiens c AK022067	XG9975	1.17
secreted protein of unknown function	N25915	1.17
nuclear receptor subfamily 6, group A, member 1	BC000738	1.17
CUG triplet repeat, RNA binding protein 1	AL120375	1.17
emelin (Emery-Dreifuss muscular dystrophy)	NM_024984	1.17
Consensus includes gb:AL120375 / FEA=EST / DB_XREF=gi:5926274 / DB_XREF=est:DKFZp761C088_s1 / CLONE=DKFZp761C088 / JG=Hs.18653 ESTs	NM_005232	1.17
hypothetical protein FLJ12193	NM_002780	1.17
EphA1	L31782	1.17
pregnancy specific beta-1-glycoprotein 4	NM_014221	1.17
carcinoembryonic antigen-related cell adhesion molecule 7	BC001152	1.17
mature T-cell proliferation 1	NM_016025	1.17
growth arrest-specific 7	AB023430	1.17
CGI-81 protein	Z77851	1.17
protein tyrosine phosphatase, non-receptor type substrate 1	NM_005454	1.17
small inducible cytokine subfamily A (Cys-Cys), member 13	NM_018207	1.17
cereus 1 homolog, cysteine knot superfamily (Xenopus laevis)	NM_014122	1.17
hypothetical protein FLJ10739	NM_006862	1.17
Consensus includes gb:Z21819 / DEF=H. sapiens carcinoembryonic antigen gene / FEA=mRNA / DB_XREF=gi:437783 / JG=Hs.166040 H. sapiens carcinoembryonic antigen Z21819	AF103530	1.17
PRO0245 protein	AF063591	1.17
tudor and KH domain-containing protein	NM_002920	1.17
immunoglobulin kappa constant	BE258133	1.17
antigen identified by monoclonal antibody MRC OX-2		
regulatory factor X, 4 (influences HLA class II expression)		
chromobox homolog 2 (Pc class homolog, Orsoptilia)		

angiotensin 1	NM_001146	1.17
cytochrome P450, subfamily 1 (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile)	AU144855	1.17
WD repeat domain 3	NM_006784	1.17
Z'-5'-oligoadenylate synthetase 3 (100 kD)	NM_006187	1.17
annexin A13	NM_004306	1.17
RAS guanyl releasing protein 2 (calcium and DAG-regulated)	AI688812	1.17
hypothetical protein FLJ21347	NM_022827	1.17
mucin 1, transmembrane	NM_002456	1.17
hypothetical protein FLJ12994	NM_022841	1.17
carbonic anhydrase XII	BC000278	1.17
T-cell acute lymphocytic leukemia 1	X51990	1.17
gb:AB058895.1 /DEF=Homo sapiens GHRHRps mRNA for growth hormone releasing hormone receptor, pituitary splice variant, complete cds. /FEA=mRNA /GEN=GHRH /AB058895		1.17
similar to proline-rich protein 48		1.17
KIAA1624 protein	NM_019043	1.17
Consensus includes gb:AA865601 /FEA=EST /DB_XREF=est093003.s1 /CLONE=IMAGE:1455868 /UG=Hs.128281 Homo sapiens Chromosome	NM_020860	1.17
a disintegrin and metalloproteinase domain 30	NM_021794	1.17
casein, beta	NM_001891	1.17
hypothetical protein FLJ20435	NM_017821	1.17
solute carrier family 29 (nucleoside transporters), member 1	NM_004955	1.17
soluble adenylyl cyclase	AL035122	1.17
dihydropyrimidine dehydrogenase	NM_000110	1.17
lysosomal-associated protein transmembrane 4 alpha	NM_014713	1.17
LIM domains containing 1	NM_014240	1.17
SH3 protein interacting with Nck 90 kDa	NM_016453	1.17
expressed in activated T/LAK lymphocytes	NM_007287	1.17
hypothetical protein FLJ22029	NM_024949	1.17
deleted in ataxia-perlimia-like	NM_001351	1.17
gb:AJ25093.1 /DEF=Homo sapiens mRNA for single-chain antibody, complete cds (scFv2). /FEA=CDS /PROD=immunoglobulin /DB_XREF=gi:3090427 /UG=Hs.159386		1.17
lucosyltransferase 3 (galactoside 3(4)-L-Lucosyltransferase, Lewis blood group included)	AW080549	1.17
CD14 antigen	NM_006591	1.17
supervillin	AF051851	1.17
adenovirus 5 E1A binding protein	BE250417	1.17
unc-13-like (C. elegans)	NM_006377	1.17
osmosis responsive factor	NM_012382	1.17
gb:NM_016528.1 /DEF=Homo sapiens hydroxyacid oxidase 3 (medium-chain) (HAO3), mRNA. /FEA=mRNA /GEN=HAO3 /PROD=hydroxyacid oxidase 3 /DB_XREF=gi:771 /NM_016528		1.17
cold autoimmune syndrome 1	NM_004895	1.17
ATPase, Class VI, type 11B	AB023173	1.17
Consensus includes gb:Y10259.1 /DEF=H sapiens ACTH receptor mRNA 3UTR. /FEA=mRNA /DB_XREF=gi:2230860 /UG=Hs.274501 H.sapiens ACTH receptor mRNA 3UTR		1.17
KIAA0844 protein	NM_014851	1.17
prophet of P11, paired-like homeodomain transcription factor	AI635302	1.17
nuclear receptor coactivator 6	AF076215	1.17
hypothetical protein	AF128458	1.17
ectodysplasin A2 isoform receptor	BC006214	1.17
splicing factor 3a, subunit 2, 66kD	NM_021783	1.17
DKFZP547E2110 protein	L21980	1.17
early growth response 2 (Krox-20 homolog, Drosophila)	BF062335	1.17
RAB22A, member RAS oncogene family	NM_000399	1.17
gonadotropin-releasing hormone 2	NM_020673	1.17
keratin 2A (epidermal Ichthyosis bullosa of Siemens)	NM_001501	1.17
GREB1 protein	NM_000423	1.17
Consensus includes gb:BE045982 /FEA=EST /DB_XREF=gi:8363035 /DB_XREF=est091b04.x4 /CLONE=IMAGE:2916847 /UG=Hs.99093 Homo sapiens chromosome	AF245390	1.17
hypothetical protein FLJ21988	NM_022493	1.17
chromosome 8 open reading frame 1	AI738702	1.17
ATP-binding cassette, sub-family C (CFTR/MRP), member 6	AI084637	1.17
protocadherin alpha 3	AF152481	1.17
potassium inwardly-rectifying channel, subfamily J, member 4	NM_004981	1.17
DKFZP134D193 protein	AK000323	1.17
hypothetical protein	NM_016623	1.17
hypothetical protein MGC16714	BE782298	1.17
potassium voltage-gated channel, Isk-related family, member 2	NM_005136	1.17
inactivation escape 1	NM_003869	1.17
Consensus includes gb:AK024135.1 /DEF=Homo sapiens cDNA FLJ14073 fls, clone HEMBB1001812. /FEA=mRNA /DB_XREF=gi:10436441 /UG=Hs.296751 Homo sapie	AK024135	1.17

proteoglycan 2, bone marrow (natural killer cell activator, eosinophil granule major basic protein) 1.17 BC005929
 ATP-binding cassette, sub-family C (CFTR/MRP), member 9 1.17 NM_020297
 C-type lectin-like receptor-1 1.17 NM_016511
 a disintegrin and metalloproteinase domain 7 1.17 AF215824
 zinc finger protein 281 1.17 NM_005086
 calmodulin 3 (phosphorylase kinase, delta) 1.17 AV65208
 H4 histone family, member B 1.17 NM_003539
 Consensus includes gb:A824954 /FEA=EST /DB_XREF=gi:5445625 /DB_XREF=est:w004c07.x1 /CLONE=IMAGE:2304684 /UG=Hs.157429 SRY (sex determining region) 1.17 AB24954
 truncated repeat containing 11 (THR-associated protein, 230 kDa subunit) 1.17 NM_005120
 RAB2, member RAS oncogene family-like 1.17 AL050259
 solute carrier family 16 (monocarboxylic acid transporters), member 5 1.17 NM_004663
 bromodomain adjacent to zinc finger domain, 1A 1.17 NM_004685
 small inducible cytokine subfamily A (Cys-Cys), member 24 1.17 NM_013448
 DKFZP58H2123 protein 1.17 NM_002991
 RBP2 interacting protein 8 1.17 AB71186
 Consensus includes gb:AK025271.1 /DEF=Homo sapiens cDNA: FLJ21618 fls, clone COL07487, /FEA=mRNA /DB_XREF=gi:10437753 /UG=Hs.306790 Homo sapiens cDNA: AK025271 1.17 AI197870
 hypothetical protein FLJ23451 1.17 NM_024766
 Consensus includes gb:AL049430.1 /DEF=Homo sapiens mRNA: cDNA DKFZb586H201 (from clone DKFZp586H201), /FEA=mRNA /DB_XREF=gi:4500210 /UG=Hs.3020 AL049430 1.17 U47824
 dentatorubral-pallidolysian atrophy (atrophin-1) 1.17 AF152505
 protocadherin gamma subfamily A, 11 1.17 AL137316
 KIAA1609 protein 1.17 NM_001841
 cannabinoid receptor 2 (macrophage) 1.17 BC005338
 Consensus includes gb:AK024938.1 /DEF=Homo sapiens cDNA: FLJ21285 fls, clone COL01912, /FEA=mRNA /DB_XREF=gi:10437359 /UG=Hs.287655 Homo sapiens cDNA: AK024938 1.17 AK024938
 hypothetical protein LOC37147 1.17 NM_070423
 retinol X receptor, alpha 1.17 BE675800
 SH2 domain protein 2A 1.17 NM_003975
 Consensus includes gb:BG434174 /FEA=EST /DB_XREF=gi:13340680 /DB_XREF=est:602506106F1 /CLONE=IMAGE:4603513 /UG=Hs.23437 Homo sapiens cDNA: FLJ1 BG434174 1.17 AF249871
 NK3 transcription factor homolog A (Drosophila) 1.17 U65065
 protein tyrosine phosphatase, receptor type, N polypeptide 2 1.17 AA121502
 KIAA1237 protein 1.17 L10038
 kallikrein 1, renal/pancreas/salivary 1.17 NM_000440
 phosphodiesterase 6A, cGMP-specific, rod, alpha 1.17 Z39557
 Consensus includes gb:Z39557 /FEA=EST /DB_XREF=gi:562749 /DB_XREF=est:HSC1DB062 /CLONE=c-10b06 /UG=Hs.12533 Homo sapiens clone 23705 mRNA seque 1.17 M31172
 transforming growth factor, alpha 1.17 NM_016508
 cyclin-dependent kinase-like 3 1.17 AA522816
 Cluster Ind. AA522816/n40e12.a1 Homo sapiens cDNA, 3' end /clone=IMAGE-979342 /clone_end=3 /gb=AA522816 /gi=2263528 /ug=Hs.173789 /len=564 1.17 AJ010070
 partial transcript encompassing THC211630 gene 1.17 BF303939
 similar to RIKEN cDNA 231004OG17 gene 1.17 AI247566
 thioredoxin reductase 2 1.17 L35901
 cholinergic receptor, nicotinic, alpha polypeptide 4 1.17 AW874481
 Consensus includes gb:AW974481 /FEA=EST /DB_XREF=gi:8165680 /DB_XREF=est:EST386585 /UG=Hs.293755 ESTs 1.17 NM_004527
 mesenchyme homeo box 1 1.17 BC002827
 topomycin 4 1.17 NM_005693
 Consensus includes gb:AL535113 /FEA=EST /DB_XREF=gi:12798606 /DB_XREF=est:AL535113 /CLONE=CS00F008YC23 (3 prime) /UG=Hs.283006 phospholipase C, b 1.17 AL131051
 KIAA1277 protein 1.17 NM_020482
 activator of CREM in testis 1.17 NM_020482
 Consensus includes gb:AK026106.1 /DEF=Homo sapiens cDNA: FLJ22453 fls, clone HRC09879, highly similar to AF059516 Homo sapiens telodiol-like 2 protein (TLL2) mR 1.17 AK026106
 Consensus includes gb:AF035290.1 /DEF=Homo sapiens clone 23556 mRNA sequence, /FEA=mRNA /DB_XREF=gi:2661044 /UG=Hs.106300 Homo sapiens clone 23556 AF035290 1.17 AF035290
 neuronal Src adaptor homolog 1.17 AV705938
 gamma-aminobutyric acid (GABA) A receptor, epsilon 1.17 NM_004961
 fasciculation and elongation protein zeta 2 (zyglin II) 1.17 AB65892
 Misshapen/Nik-related kinase 1.17 AL157418
 seizure related 6 homolog (mouse)-like 1.17 AB023144
 low density lipoprotein receptor-related protein 6 1.17 NM_002336
 gb:NM_004250.1 /DEF=Homo sapiens RAB9, member RAS oncogene family, pseudogene 1 (RAB9P1), mRNA, /FEA=mRNA /GEN=RAB9P1 /PROD=RAB9, member RAS 1.17 NM_004250
 butyrophilin, subfamily 3, member A1 1.17 NM_013233
 Shc-20 related kinase 1.17 NM_002170
 interferon, alpha 6 1.17 BC005563
 solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2 1.17
 Consensus includes gb:AA543076 /FEA=EST /DB_XREF=gi:2291556 /DB_XREF=est:n96102.s1 /CLONE=IMAGE:927771 /UG=Hs.82171 Homo sapiens (clone 115392) m 1.17 AA543076

Consensus includes gb:AA716107 /FEA=EST /DB_XREF=gi:2728381 /DB_XREF=est:zg3d06.s1 /CLONE=IMAGE:398027 /UG=Hs.8963 Homo sapiens mRNA full length
 tropomyosin 1 (alpha) NM_000386 1.17
 sperm associated antigen 1 AF311312 1.17
 collagen, type XV, alpha 1 NM_001855 1.17
 phosphatidylinositol glycan, class B AU144243 1.17
 histidine decarboxylase NM_002112 1.17
 Consensus includes gb:AW879289 /FEA=EST /DB_XREF=gi:8170577 /DB_XREF=est:EST391399 /UG=Hs.293472 ESTs
 gb:M73068.1 /DEF=Human androgen receptor mutant gene, mRNA, complete cds. /FEA=mRNA /GEN=AR /PROD=androgen receptor /DB_XREF=gi:178655 /FL=gb:M730
 hypoxia-inducible factor 1, alpha subunit inhibitor AW879289 1.16
 secreted frizzled-related protein 1 NM_017902 1.16
 hypothetical protein FLJ21820 NM_003012 1.16
 putative non-coding transcript (DiGeorge critical region 5) AK021672 1.16
 hypothetical protein FLJ20516 A479306 1.16
 hypothetical protein FLJ23459 NM_017858 1.16
 chromodomain helicase DNA binding protein 3 NM_024775 1.16
 a disintegrin and metalloproteinase domain 8 BE378542 1.16
 chromosome 11 open reading frame 16 NM_001109 1.16
 myosin X NM_020643 1.16
 cysteine sulfonic acid decarboxylase-related protein 2 A1581354 1.16
 calponin like transmembrane domain protein NM_015989 1.16
 Consensus includes gb:BF224076 /FEA=EST /DB_XREF=gi:11131303 /DB_XREF=est:7q83a08.x1 /CLONE=IMAGE:3704942 /UG=Hs.307438 Human DNA sequence from
 DnaJ (Hsp40) homolog, subfamily B, member 12 NM_024734 1.16
 protein phosphatase 1A (lomerty 2C), magnesium-dependent, alpha isoform BG283782 1.16
 protocadherin beta 3 NM_021003 1.16
 U5 snRNP-specific protein, 200 kDa (DEXH RNA helicase family) NM_018937 1.16
 Consensus includes gb:AA476303 /FEA=EST /DB_XREF=gi:2204514 /DB_XREF=est:zw29d09.s1 /CLONE=IMAGE:770705 /UG=Hs.306357 Homo sapiens isolate donor N
 hypothetical protein KIAA0397 gene product Z70200 1.16
 gamma-aminobutyric acid (GABA) receptor, rho 2 NM_014306 1.16
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily I, member 1 NM_002043 1.16
 decortin NM_006015 1.16
 KIAA0669 gene product A1281593 1.16
 phosphatidylinositol 4-kinase, catalytic, beta polypeptide NM_030797 1.16
 gap junction protein, beta 3, 31kD (connexin 31) AB007857 1.16
 olfactory receptor, family 1, subfamily F, member 2 NM_005221 1.16
 Consensus includes gb:X07868 /DEF=Human DNA for insulin-like growth factor II (IGF-2); exon 7 and additional ORF /FEA=mRNA_1 /DB_XREF=gi:32998 /UG=Hs.25166
 transient receptor potential channel 1 A912351 1.16
 KIAA0669 gene product Z73803 1.16
 phosphatidylinositol 4-kinase, catalytic, beta polypeptide AF201292 1.16
 gap junction protein, beta 3, 31kD (connexin 31) NM_002651 1.16
 olfactory receptor, family 1, subfamily F, member 2 NM_024928 1.16
 hypothetical protein FLJ10759 AU003145 1.16
 Thy-1 cell surface antigen R71157 1.16
 septin 6 AL558479 1.16
 Sec23 homolog A (S. cerevisiae) D50918 1.16
 hepatocellular carcinoma-associated gene TD28 A173659 1.16
 HCG11-7 protein NM_018687 1.16
 deleted in colorectal carcinoma X81001 1.16
 interferon induced transmembrane protein 2 (1-8D) NM_005215 1.16
 vesicle-associated membrane protein 1 (synaptobrevin 1) NM_006435 1.16
 anti-Müllerian hormone AU150319 1.16
 hypothetical protein FLJ13150 NM_004479 1.16
 putative Rab5-interacting protein NM_024813 1.16
 MHC class I polypeptide-related sequence A AL050318 1.16
 MAWD binding protein NM_000247 1.16
 mitogen-activated protein kinase kinase 12 NM_022129 1.16
 neurogenic differentiation 1 AW025150 1.16
 autophagy inducer (autophagy inducer) NM_024112 1.16
 decay accelerating factor for complement (CD55, Cr1, blood group system) NM_002500 1.16
 carbon-chloride cotransporter-interacting protein NM_006558 1.16
 KIAA0197 protein NM_000574 1.16
 epilepsy, progressive myoclonus type 2, Lafora disease (laforin) AK026238 1.16
 NM_005870 1.16

testis expressed sequence 15	NM_031271	1.16
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)	AI73248	1.16
ribosomal protein L3-like	NM_005061	1.16
cadherin 6, type 2, K-cadherin (fetal kidney)	NM_004932	1.16
KIAA1548 protein	NM_020909	1.16
scf-Coenzyme A oxidase 1, palmityl	S69189	1.16
DV527-related protein	AB024518	1.16
Consensus includes gb:AK027217.1 /DEF=Homo sapiens cDNA: FLJ23564 fls, clone LNG10773. /FEA=mRNA /DB_ XREF=gi:10440296 /UG=Hs.306913 Homo sapiens cC AK027217	AB024518	1.16
lipase protein	AF225418	1.16
hypothetical protein FLJ12177	NM_025033	1.16
KIAA1701 protein	H11725	1.16
NCK adaptor protein 2	BC000103	1.16
olfactory receptor, family 7, subfamily A, member 17	AC005255	1.16
KIAA0429 gene product	AI027678	1.16
KIAA1007 protein	NM_016284	1.16
signal transducer and activator of transcription 5A	NM_003152	1.16
Consensus includes gb:BE646315 /FEA=EST /DB_ XREF=est:764108.x1 /CLONE=IMAGE.3289191 /UG=Hs.292503 ESTs, Weakly similar to T471 BE646315	NM_003152	1.16
nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)	BC002844	1.16
Consensus includes gb:AK025064.1 /DEF=Homo sapiens cDNA: FLJ21411 fls, clone COL03986. /FEA=mRNA /DB_ XREF=gi:10437503 /UG=Hs.306758 Homo sapiens cC AK025064	BC002844	1.16
Consensus includes gb:AK025363.1 /DEF=Homo sapiens cDNA: FLJ21710 fls, clone COL10087. /FEA=mRNA /DB_ XREF=gi:10437865 /UG=Hs.306807 Homo sapiens cC AK025363	AI346187	1.16
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	NM_001713	1.16
betaine-homocysteine methyltransferase	NM_003318	1.16
TTK protein kinase	AL503687	1.16
Consensus includes gb:AL583687 /FEA=EST /DB_ XREF=gi:12952889 /DB_ XREF=est:AL583687 /CLONE=CSDDJ008YL09 (5 prime) /UG=Hs.134852 ESTs	NM_005711	1.16
EGF-like repeats and discoidin I-like domains 3	AL521391	1.16
ELL gene (11-19 lysine-rich leukemia gene)	NM_020806	1.16
gephyrin	M65217	1.16
heat shock transcription factor 2	NM_018952	1.16
homoio box B6	AK022014	1.16
hypothetical protein FLJ11852	AA114186	1.16
Consensus includes gb:AA114186 /FEA=EST /DB_ XREF=gi:1668052 /DB_ XREF=est:zm75d3.s1 /CLONE=IMAGE.564004 /UG=Hs.23964 sin3-associated polypeptide, 18 AA114186	NM_014796	1.16
KIAA0748 gene product	NM_015644	1.16
DKFZP434B103 protein	AL079292	1.16
Consensus includes gb:AK026682.1 /DEF=Homo sapiens cDNA: FLJ23029 fls, clone LNG01883. /FEA=mRNA /DB_ XREF=gi:10435989 /UG=Hs.306864 Homo sapiens cC AK026682	NM_018584	1.16
hypothetical protein	AI692203	1.16
hypothetical protein PRO1489	NM_016642	1.16
ring finger protein 3	NM_015577	1.16
spectrin, beta, non-erythrocytic 5	AF057557	1.16
Consensus includes gb:AV761014 /FEA=EST /DB_ XREF=gi:10918862 /DB_ XREF=est:AV761014 /CLONE=MDSCCF09 /UG=Hs.291972 ESTs, Moderately similar to SC14 AV761014	AF057557	1.16
retinoic acid induced 14	U44378	1.16
regulator of Fas-induced apoptosis	NM_000103	1.16
MAD, mothers against decapentaplegic homolog 4 (Drosophila)	NM_030931	1.16
cyclochrome P450, subfamily XIX (aromatization of androgens)	AB004853	1.16
epididymal secretory protein ESP13.2	U66495	1.16
chromosome 21 open reading frame 18	NM_012107	1.16
leptin receptor	BF438056	1.16
bronchodomain containing protein 75 kDa human homolog	AF228413	1.16
cAMP responsive element binding protein-like 2	NM_024049	1.16
nuclear receptor subfamily 5, group A, member 2	AK021576	1.16
hypothetical protein MGC5566	AK027071	1.16
Consensus includes gb:AK021578.1 /DEF=Homo sapiens cDNA FLJ11514 fls, clone HEMBA1002229. /FEA=mRNA /DB_ XREF=gi:10432783 /UG=Hs.306602 Homo sapie AK021576	NM_004994	1.16
transforming growth factor beta-stimulated protein TSC-22	NM_004178	1.16
matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase)	BC004155	1.16
Consensus includes gb:J04178.1 /DEF=Human abnormal beta-hexosaminidase alpha chain (HEXA) mRNA, partial cds. /FEA=mRNA /DB_ XREF=gi:184001 /UG=Hs.1862 J04178	NM_015320	1.16
ring finger protein 5	AA528252	1.16
Rho guanine nucleotide exchange factor (GEF) 4	NM_014777	1.16
KIAA0286 gene product	AF237813	1.16
KIAA0133 gene product	AB037854	1.16
NPD009 protein	NM_024541	1.16
hypothetical protein DKFZp41A023	M58051	1.16
hypothetical protein FLJ13114	NM_001976	1.16
fibroblast growth factor receptor 3 (echondroplasia, thanatophoric dwarfism)		
endolase 3, (beta, muscle)		

gb:225427.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete CDS, /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405732 /FL=gb:225427.1
 claudin 1 NM_021101 1.16
 galactosylceramidase (Krabbe disease) D25284 1.16
 GA binding protein transcription factor, beta subunit 1 (53kD) NM_005254 1.16
 CSE1 chromosome segregation 1-like (yeast) AF053841 1.16
 hypothetical protein FLJ13725 NM_024519 1.16
 KIAA0892 protein AW157731 1.16
 hypothetical protein FLJ20216 NM_017715 1.16
 myelin transcription factor 1 M86980 1.16
 protein tyrosine phosphatase, receptor type, A BF740139 1.16
 nuclear receptor binding protein NM_013392 1.16
 Epha3 AF213460 1.16
 peptidylglycine alpha-amidating monooxygenase NM_000919 1.16
 carboxypeptidase D BE349147 1.16
 signal recognition particle 54kD NM_003136 1.16
 imprinted in Prader-Willi syndrome AI672541 1.16
 RAN binding protein 8 NM_006390 1.16
 musashi homolog 1 (Drosophila) NM_002442 1.16
 tumor necrosis factor receptor superfamily, member 4 AJ277151 1.16
 dentin, light polypeptide (Lca) NM_001833 1.16
 solute carrier family 12 (potassium/chloride transporters), member 4 U55054 1.16
 Consensus includes gb:U55987 /FEA=EST /DB_XREF=est:HSU55987 /CLONE=47507 /UG=Hs.323773 ESTs. Highly similar to NPC1_HUMAN NIE U55987 1.16
 ATP-binding cassette, sub-family A (ABC1), member 7 NM_019112 1.16
 transducin (beta)-like 2 AK026529 1.16
 2',5'-oligoadenylate synthetase 1 (40-48 kD) NM_018816 1.16
 Consensus includes gb:AL046983.1 /DEF=Homo sapiens mRNA; cDNA DKFpZ564D042, /FEA=mRNA /DB_XREF=gi:4884234 /UG=Hs.2401 AL046983 1.16
 hypothetical protein FLJ20051 NM_019067 1.16
 Cluster ind. A1248730q440b01.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:1871401 /clone_end=3 /gb=A1246730 /gi=3842127 /ug=Hs.126901 /len=770 A1246730 1.16
 hypothetical protein A-211C8.1 BE363444 1.16
 myosin, light polypeptide 5, regulatory NM_002477 1.16
 spilloproteins B48 receptor NM_018690 1.16
 monocyte to macrophage differentiation-associated NM_012328 1.16
 reelin NM_005045 1.16
 ELK4, ETS-domain protein (SRF accessory protein 1) NM_021795 1.16
 cholecystokinin B receptor BC000740 1.16
 B7 protein NM_008892 1.16
 lectin, galactoside-binding, soluble, 8 (galectin 8) AF342815 1.16
 EGF-TM7-180philin-related protein NM_022159 1.16
 N-sulfoglucosamine sulfohydrolase (sulfamidase) NM_000199 1.16
 coagulation factor II (thrombin) receptor-like 2 NM_004101 1.16
 nuclear receptor subfamily 1, group H, member 2 NM_007121 1.16
 brylin-like NM_004053 1.16
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 AA772023 1.16
 KIAA1031 protein NM_025013 1.16
 interleukin 22 NM_020525 1.16
 hypothetical protein FLJ21868 NM_022769 1.16
 lung cancer candidate NM_002725 1.16
 glutathione S-transferase M4 NM_000850 1.16
 olfactory receptor, family 7, subfamily C, member 2 NM_012377 1.16
 hypothetical protein FLJ20359 NM_017781 1.16
 putative DNA binding protein BG033764 1.16
 FYN binding protein (FYB-120/130) AF198052 1.16
 piccolo (presynaptic cytomatrix protein) BC001304 1.16
 RAB7, member RAS oncogene family-like 1 BG338251 1.16
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1 U08015 1.16
 islet cell autoantigen 1 (89kD) NM_022307 1.16
 macrophage scavenger receptor 1 NM_002445 1.16
 chromosome 19 open reading frame 3 NM_005716 1.16
 fibronectin 1 X02761 1.16
 NY-REN-24 antigen BC004282 1.16
 KIAA0040 gene product NM_014656 1.16
 solute carrier family 22 (organic anion transporter), member 7 AF210455 1.16
 latent transforming growth factor beta binding protein 2 NM_000428 1.16

LIM homeobox protein 2 U11701 1.16
 synovial sarcoma translocation, chromosome 18 AF257501 1.16
 Consensus includes gb:BE409594 /FEA=EST /DB_XREF=gi:9346444 /DB_XREF=est:601299404F1 /CLONE=IMAGE:3629592 /UG=Hs.326464 Homo sapiens cDNA: FLJ: BE409594 1.16
 immunoglobulin superfamily, member 3 AB007935 1.16
 G protein-coupled receptor 75 NM_006794 1.16
 3-hydroxytryptophan 3,4-dioxygenase NM_012205 1.16
 Consensus includes gb:H94842 /FEA=EST /DB_XREF=gi:1102475 /DB_XREF=est:y57a03.31 /CLONE=IMAGE:230188 /UG=Hs.7882 ESTs H94842 1.16
 phospholipase A2, group IVA (cytosolic, calcium-dependent) M88874 1.16
 splicing factor 3b, subunit 3, 130kD O13642 1.16
 hypothetical protein FLJ11939 AK25402 1.16
 KISS-1 metastasis-suppressor NM_002256 1.16
 tropotin 1, skeletal, slow NM_003281 1.16
 KIAA0268 gene product NM_021645 1.16
 rTS beta protein NM_017512 1.16
 Consensus includes gb:AL049381.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586J2118 (from clone DKFZp586J2118). /FEA=mRNA /DB_XREF=gi:4500168 /UG=Hs.218 AL049381 1.16
 VENT-like homeobox 2 NM_014468 1.16
 hypothetical protein FLJ10346 NM_018065 1.16
 p1m-1 oncogene M24779 1.16
 importin 13 NM_014652 1.16
 plasma glutamate carboxypeptidase NM_016134 1.16
 Consensus includes gb:BF111846 /FEA=EST /DB_XREF=gi:10941536 /DB_XREF=est:7036001.x1 /CLONE=IMAGE:3523608 /UG=Hs.106334 Homo sapiens clone 23636 NM_018018 1.16
 hypothetical protein FLJ10193 BF111846 1.16
 Consensus includes gb:AK026980.1 /DEF=Homo sapiens cDNA: FLJ23327 fls, clone HEP12630, highly similar to HSNF37 Homo sapiens ZNF37A mRNA for zinc finger protein FLJ23327 NM_024574 1.16
 HTGN29 protein NM_020199 1.16
 polymerase (DNA directed), theta NM_014125 1.16
 neuropeptide FF 1; RFamide-related peptide receptor NM_022146 1.16
 mitogen-activated protein kinase kinase 4 NM_006724 1.16
 ovarian cancer overexpressed 1 NM_015945 1.16
 ATP-binding cassette, sub-family A (ABC1), member 2 AF327657 1.16
 ethandiamine kinase NM_018638 1.16
 ligand of neuronal nitric oxide synthase with carboxyl-terminal PDZ domain AF037070 1.16
 amiloride-sensitive cation channel 1, neuronal (degenerin) NM_001094 1.16
 Consensus includes gb:H53689 /FEA=EST /DB_XREF=gi:993836 /DB_XREF=est:y136g07.r1 /CLONE=IMAGE:236124 /UG=Hs.306239 Homo sapiens clone ASPBL54 in H53689 NM_013358 1.16
 peptidylarginine deiminase type 1 AK023188 1.16
 protein phosphatase 1, regulatory (inhibitor) subunit 13B NM_024920 1.16
 hypothetical protein FLJ14281 AK023188 1.16
 interleukin 3 receptor, alpha NM_024920 1.16
 KIAA0853 protein, B7-like protein AF199028 1.16
 soluble carrier family 8 (neurotransmitter transporter, betaine/GABA), member 12 U27699 1.16
 leucine zipper, putative tumor suppressor 1 N21184 1.16
 interleukin 13 receptor, alpha 1 NM_001560 1.16
 hypothetical protein FLJ23071 NM_025192 1.16
 v-jun sarcoma virus 17 oncogene homolog (avian) NM_002228 1.16
 adenylate cyclase activating polypeptide 1 (pituitary) NM_001117 1.16
 chemokine (C-C motif) receptor 7 NM_001838 1.16
 T cell receptor alpha locus AE000859 1.16
 Consensus includes gb:AL182040.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434N199 (from clone DKFZp434N199). /FEA=mRNA /DB_XREF=gi:7328079 /UG=Hs.3064 AL182040 1.16
 BCL2-like 11 (apoptosis facilitator) NM_008538 1.16
 AD024 protein AF225418 1.16
 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 U192838 1.16
 stanniocalcin 1 U48768 1.16
 cleavage and polyadenylation specific factor 6, 68kD subunit NM_007007 1.16
 lymphocyte antigen 84 homolog, activating NK-receptor; NK-p46, (mouse) AJ008121 1.16
 GLI-Kruppel family member GLI3 (Grag cephalopolysyndactyly syndrome) NM_000168 1.16
 KIAA0014 gene product NM_014685 1.16
 Consensus includes gb:BC004327.1 /DEF=Homo sapiens, clone IMAGE:3629896. mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3629896) /DB_XI BC004327 1.16
 KIAA1841 protein NM_025190 1.16
 zinc finger protein 144 (Met-18) NM_007144 1.16
 Consensus includes gb:AK022094.1 /DEF=Homo sapiens cDNA FLJ12032 fls, clone HEMBB1001880. /FEA=mRNA /DB_XREF=gi:10433415 /UG=Hs.151194 Homo sapiens clone AK022094 1.16
 phospholipase A2, group IIE NM_014589 1.16
 Consensus includes gb:AL109707.1 /DEF=Homo sapiens mRNA full length Insert cDNA clone EUROMIMAGE 295344. /FEA=mRNA /DB_XREF=gi:5889824 /UG=Hs.943271 AL109707 1.16
 Consensus includes gb:L41657 /DEF=Homo sapiens synthetic myelin basic protein 21.5 kDa isoform gene, complete cds /FEA=CDS /DB_XREF=gi:1162921 /UG=Hs.2480 L41657 1.16

v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian)		
guanylate cyclase activator 1A (refine)	AF177761	1.16
HMB-Inducible	L36861	1.16
protein tyrosine phosphatase type IVA, member 1	AW665096	1.16
mitogen-activated protein kinase 14	AL578310	1.16
phorodin-like protein MDS019	L35253	1.16
rbp3 GTPase-activating protein, non-catalytic subunit (150kD)	BF508948	1.16
leukocyte tyrosine kinase	BF240652	1.16
growth arrest-specific 7	X52213	1.16
hypothetical protein FLJ20156	BE439987	1.16
hypothetical protein FKSG28	NM_017691	1.16
KIAA0382 protein	NM_030929	1.16
laminin, beta 2 (laminin S)	A1348094	1.16
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)	X79683	1.16
hypothetical protein FLJ20244	NM_003250	1.16
actin-like 6	AL390133	1.16
flightless I homolog (Drosophila)	NM_016188	1.16
tumor protein D52	U80184	1.16
hypothetical protein FLJ13159	AA524023	1.16
KIAA0082 protein	NM_021940	1.16
CD163 antigen	D43949	1.16
LPS-induced TNF-alpha factor	Z22970	1.16
v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	AB034747	1.16
hypothetical protein MGC11352	W80678	1.16
S100 calcium binding protein A2	NM_030927	1.16
hypothetical protein DKFZp564F013	NM_005978	1.16
retinoic acid receptor, gamma	AL050389	1.16
CGI-01 protein	M57707	1.16
KIAA0013 gene product	AK001172	1.16
glutaredoxin 2	NM_014783	1.16
crystallin, zeta (guinone reductase)	NM_016066	1.16
chromosome 9 open reading frame 9	NM_001889	1.16
Consensus includes gb:AK000244.1 /DEF=Homo sapiens cDNA FLJ2037 fs, clone COLF5879, highly similar to AB023230 Homo sapiens mRNA for KIAA1013 protein, fl	NM_018956	1.15
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	M61213	1.15
leukocyte receptor cluster (LRC) member 5	NM_002010	1.15
LCA1-like lysophospholipase	NM_024075	1.15
Consensus includes gb:AI446234 /FEA=EST /DB_XREF=gi:4293782 /DB_XREF=est:j25g10.x1 /CLONE=IMAGE:2142594 /UG=Hs.156072 Human pre-TNK cell associate	AL110209	1.15
dibutyrate, iodotyrosine, type II	AB041844	1.15
spodin 1, (I-spondin) extracellular matrix protein	AB016305	1.15
hypothetical protein DKFZp34L0850	NM_017558	1.15
potassium inwardly-rectifying channel, subfamily J, member 6	N39314	1.15
zinc finger protein 73 (Cos12)	NM_012480	1.15
ubiquitin specific protease 1	AW489935	1.15
glycoprotein 2 (zymogen granule membrane)	NM_016295	1.15
T cell receptor gamma constant 2	AL580863	1.15
hypothetical protein DKFZp34H2215	NM_017559	1.15
CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)	NM_006889	1.15
tumor necrosis factor, alpha-induced protein 1 (endothelial)	BC001643	1.15
KIAA1451 protein	AL049923	1.15
matrix metalloproteinase 11 (stromelysin 3)	NM_005840	1.15
hippocampin-like protein 4	AL136591	1.15
glycophorin B (includes Ss blood group)	U05255	1.15
forkhead box J1	NM_001454	1.15
hypothetical protein FLJ21749	NM_025124	1.15
Gardner-Rasheed feline sarcoma viral (v-igr) oncogene homolog	NM_065248	1.15
retinaldehyde binding protein 1	NM_000326	1.15
KIAA0275 gene product	A1952009	1.15
deleted in esophageal cancer 1	NM_017418	1.15
RAR (RAS like GTPASE)	BE965869	1.15
estrogen receptor 1	NM_000125	1.15
Nilemgen breakage syndrome 1 (nibrin)	A1766269	1.15
CD2 antigen (p50), sheep red blood cell receptor	NM_001767	1.15

Consensus includes gb:AW675473 /FEA=EST /DB_XREF=gi:7540699 /DB_XREF=est:ba49g05.x1 /CLONE=IMAGE:2899928 /UG=Hs.101916 Homo sapiens mRNA; cDN / AW675473
CDC14 cell division cycle 14 homolog A (S. cerevisiae)
daudin 3
hypothetical protein FLJ22378
chloride intracellular channel 3
ATP-binding cassette, sub-family B (MDR/TAP), member 1
apolipoprotein E
hypothetical protein FLJ21032
Carbonic anhydrase-related protein 10
lymphocyte antigen 117
transient receptor potential channel 5
brain and nosopharyngeal carcinoma susceptibility protein
Consensus includes gb:AU148274 /FEA=EST /DB_XREF=gi:11009795 /DB_XREF=est:AU148274 /CLONE=MAMMA1003089 /UG=Hs.301201 Homo sapiens cDNA FLJ14148274
putative transmembrane protein
polycystic kidney disease 2-like 2
sulfolipase family, cytosolic, 1C, member 1
lymphocyte antigen 95 (activating NK-receptor; NK-p44)
cytochrome P450, subfamily IIIA, polypeptide 7
NY-REN-24 antigen
zinc finger protein
putative receptor (family A group 5)
aquaporin 4
natural killer cell receptor 2B4
CD38 antigen (collagen type I receptor, thrombospondin receptor)
small membrane protein 1
hypothetical protein FLJ14360
nuclear RNA export factor 2
MAX protein
kallikrein 11
gb:BC006370.1 /DEF=Homo sapiens. Similar to Inversin, clone MGC:12888, mRNA, complete cds. /FEA=mRNA /PROD=Similar to Inversin /DB_XREF=gi:13623522 /FL=BC006370
doublecortin; lissencephaly, X-linked (doublecortin)
polymerase (RNA) II (DNA directed) polypeptide A (220kD)
androgen receptor-associated coregulator 267
neurotrophin
Consensus includes gb:AU160041 /FEA=EST /DB_XREF=gi:11021562 /DB_XREF=est:AU160041 /CLONE=Y79AA1000989 /UG=Hs.120858 Homo sapiens cDNA FLJ136160041
heparan sulfatase (glucosaminase) 3-O-sulfotransferase 1
HLA-B associated transcript 2
Consensus includes gb:AA72320 /FEA=EST /DB_XREF=gi:4334410 /DB_XREF=est:gb7c02.x1 /CLONE=IMAGE:2148482 /UG=Hs.48504 ESTs. Moderately similar to ALU
zinc finger protein 135 (clone pTZ-17)
kinesin family member 5C
Consensus includes gb:AL109882.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 35394. /FEA=mRNA /DB_XREF=gi:5685804 /UG=Hs.23777 H AL109882
zona pellucida glycoprotein 2 (sperm receptor)
phosphorylase kinase, beta
leukocyte receptor cluster (LRC) member 4
chromogranin B (secretogranin 1)
Dystonia musculorum of mouse, human homolog of
histone H2A.F/Z variant
prostate androgen-regulated transcript 1
serine dehydratase
collagen, type IX, alpha 2
Interleukin 13 receptor, alpha 1
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 duncal homolog, Drosophila)
mitogen-activated protein kinase kinase 6
interleukin 8
Consensus includes gb:AV763520 /FEA=EST /DB_XREF=gi:10921368 /DB_XREF=est:AV763520 /CLONE=MDSCH03 /UG=Hs.282113 ESTs
cadherin 5, type 2 (T1-cadherin)
nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)
KIAA1139 protein
KIAA1045 protein
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F
DKFZP566C0424 protein
putative integral membrane transporter
Consensus includes gb:AL137437.1 /DEF=Homo sapiens mRNA; cDNA DKFZP566G231.1 (from clone DKFZP566G231). /FEA=mRNA /DB_XREF=gi:6808035 /UG=Hs.2723 AL137437

F-box only protein 5	NM_012177	1.15
Cluster Inci. AB06793:wf15405.x1 Homo sapiens cDNA, 3' end (clone IMAGE-2350865 (clone_end=3 (gb=AB06793 (gi=5393359 (ug=Hs.239450 (len=555	AB06793	1.15
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	NM_000267	1.15
hypothetical protein FLJ20375	NM_017794	1.15
interleukin 15 receptor, alpha	NM_002189	1.15
hypothetical protein FLJ11274	NM_016375	1.15
spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxin 1)	AW235612	1.15
KIAA1055 protein	AL137303	1.15
hypothetical protein FLJ23506	NM_024833	1.15
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9	MZ1940	1.15
insulin receptor	AA485908	1.15
hypothetical protein MQC3560	NM_019054	1.15
SRY (sex determining region Y-box 5	NM_006940	1.15
claudin 11 (oligodendrocyte transmembrane protein)	AF068863	1.15
hypothetical protein FLJ12387 similar to kinesin light chain	NM_022822	1.15
hypothetical protein FLJ20309	NM_017759	1.15
homeodomain interacting protein kinase 3	AF305239	1.15
hypothetical protein PRO2949	NM_018544	1.15
aldehyde dehydrogenase 1 family, member B1	NM_000692	1.15
Consensus includes gb:AB041269.1 /DEF=Homo sapiens mRNA for keratin 19, partial cds, isolate K19-141 /FEA=mRNA /GEN=keratin 19 /PROD=keratin 19 /DB_XREF=AB041269	NM_000692	1.15
ankyrin 2, neuronal	BF728212	1.15
hypothetical protein FLJ13511	AF209931	1.15
KIAA1037 protein	AK023778	1.15
H1 histone family, member 4	NM_005321	1.15
protein phosphatase 2, regulatory subunit B (B56), beta isoform	NM_006244	1.15
CCAAT-box-binding transcription factor	NM_005760	1.15
UDP-N-acetylglucosamine-2-epimerase/N-acetylmannosamine kinase	NM_005476	1.15
adenylylase kinase 5	NM_012093	1.15
prion protein (PrP ^{Sc}) (Creutzfeldt-Jakob disease, Gerstmann-Sträussler-Scheinker syndrome, fatal familial insomnia)	AV725328	1.15
sterc-C5-desaturase (fungal ERG3, delta-5-desaturase)-like	D85181	1.15
small inducible cytokine subfamily A (Cys-Cys), member 21	NM_002989	1.15
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 5	NM_006840	1.15
signal transducer and activator of transcription 1, 91KD	M97935	1.15
hypothetical protein FLJ11160	NM_016344	1.15
TATA box binding protein (TBP)-associated factor, RNA polymerase II, S, 150KD	NM_006706	1.15
adipose most abundant gene transcript 1	NM_004787	1.15
KIAA0728 gene product	NM_014718	1.15
phosphatidylethanolamine N-methyltransferase	NM_007169	1.15
stromal cell-derived factor 1	U18495	1.15
triple helix repeat containing 21	U80756	1.15
spondin 2, extracellular matrix protein	NM_012445	1.15
sulfite oxidase	AA129776	1.15
HSPC054 protein	NM_014152	1.15
KIAA0391 gene product	AA129755	1.15
KIAA1028 protein	A1738802	1.15
putative Ly-6 superfamily member	NM_024123	1.15
hypothetical protein PRO0889	NM_018496	1.15
solute carrier family 21 (organic anion transporter), member 11	NM_013272	1.15
Consensus includes gb:AL390149.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547B026 (from clone DKFZp547B026) /FEA=mRNA /DB_XREF=gi:3368883 (UG=Hs.2638) /AL390149	NM_013272	1.15
a diisolegmin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif 6	NM_014273	1.15
glutaminase	AF097495	1.15
STAT induced STAT inhibitor-4	NM_004232	1.15
calcium/calmodulin-dependent protein kinase IG	AL049688	1.15
egl-like module containing, much-like, hormone receptor-like sequence 2	NM_013447	1.15
synaptonemal complex protein 1	NM_003176	1.15
proteoglycan gamma subfamily B, 5	AF152521	1.15
regulator of G-protein signalling 4	AL514445	1.15
Consensus includes gb:AF070591.1 /DEF=Homo sapiens clone 24649 mRNA sequence /FEA=mRNA /DB_XREF=gi:3387968 (UG=Hs.322473 Homo sapiens clone 24649	NM_001908	1.15
cathelin B	NM_001310	1.15
CAMP responsive element binding protein-like 2	AF78576	1.15
RE1-silencing transcription factor	U63542	1.15
Consensus includes gb:U63542.1 /DEF=Human putative FAP protein mRNA, partial cds. /FEA=mRNA /PROD=FAP protein /DB_XREF=gi:1890646 (UG=Hs.166101 Human	U63542	1.15
KIAA1097 protein	AK023664	1.15

triple functional domain (PTPRF interacting)
 kalikrein 13
 N-acetylglucosaminidase, alpha-
 CULL8 protein
 cholinergic receptor, nicotinic, alpha polypeptide 5
 protein tyrosine phosphatase, receptor type, R
 hypothetical protein FLJ11726
 Consensus includes gb:AK026484.1 /DEF=Homo sapiens cDNA: FLJ22831 fls, clone KIAA1461. /FEA=mRNA /DB_XREF=gi:10439356 /UG=Hs.321668 Homo sapiens cDNA: AK026484
 salivary proline-rich protein
 melanoma inhibitory activity
 transforming, acidic coiled-coil containing protein 1
 peptidyl-prolyl isomerase G (cyclophilin G)
 class I cytokine receptor
 stromal antigen 2
 complement component (3d/Epstein Barr virus) receptor 2
 Consensus includes gb:AU147528 /FEA=EST /DB_XREF=gi:11009049 /DB_XREF=est:AU147528 /CLONE=MAMMA1000940 /UG=Hs.301811 Homo sapiens cDNA: FLJ14
 Consensus includes gb:AK024138.1 /DEF=Homo sapiens cDNA: FLJ14076 fls, clone HEMBB1001825. /FEA=mRNA /DB_XREF=gi:10439445 /UG=Hs.306687 Homo sapie
 tropoin 1, cardiac
 KIAA0218 gene product
 CGI-78 protein
 protein C receptor, endothelial (EPCR)
 Kruppel-like factor 7 (ubiquitous)
 ubiquitously transcribed tetratricopeptide repeat gene, X chromosome
 Consensus includes gb:AK024381.1 /DEF=Homo sapiens cDNA: FLJ14319 fls, clone PLACE3000406. /FEA=mRNA /DB_XREF=gi:10436753 /UG=Hs.287638 Homo sapie
 Consensus includes gb:BE872181 /FEA=EST /DB_XREF=gi:10032712 /DB_XREF=est:7b51c08.x1 /CLONE=IMAGE:3231758 /UG=Hs.155489 NS1-associated protein 1 /F
 interferon, gamma-inducible protein 16
 prostaglandin E receptor 3 (subtype EP3)
 SGC32445 protein
 core-binding factor, runt domain, alpha subunit 2; translocated to 1; cyclin D-related
 potassium voltage-gated channel, delayed-rectifier, subfamily S, member 1
 cut-like 1, CCAAT displacement protein (Drosophila)
 hypothetical protein FLJ20055
 hypothetical protein FLJ22638
 G protein-coupled receptor, family C, group 5, member B
 AUP-ribosylarginine hydrolase
 Consensus includes gb:AF070620.1 /DEF=Homo sapiens clone 24694 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283886 /UG=Hs.158550 Homo sapiens clone 24694
 thyroid hormone receptor-associated protein, 150 kDa subunit
 glycine dehydrogenase (decarboxylating; glycine decarboxylase, glycine cleavage system protein P)
 isovaleryl Coenzyme A dehydrogenase
 Consensus includes gb:AU144792 /FEA=EST /DB_XREF=gi:11006313 /DB_XREF=est:AU144792 /CLONE=HEMBA1002973 /UG=Hs.319540 Homo sapiens cDNA: FLJ10
 caspase recruitment domain family, member 15
 mannosyl (alpha-1,3)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
 G protein-coupled receptor 19
 septin 8
 hypothetical protein FLJ10063
 Consensus includes gb:AU146655 /FEA=EST /DB_XREF=gi:11008176 /DB_XREF=est:AU146655 /CLONE=HEMBA1001133 /UG=Hs.288178 Homo sapiens cDNA: FLJ11
 Consensus includes gb:BF062299 /FEA=EST /DB_XREF=gi:10821197 /DB_XREF=est:7N76c03.x1 /CLONE=IMAGE:3481325 /UG=Hs.323079 Homo sapiens mRNA; cDN
 delta-tubulin
 KIAA0657 protein
 Fc fragment of IgG, low affinity IIb, receptor for (CD32)
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14)
 microtubule-associated protein 4
 ankyrin 1, erythrocytic
 cocaine- and amphetamine-regulated transcript
 purinergic receptor P2X, ligand-gated ion channel, 2
 obaferrin
 SRY (sex determining region Y)-box 22
 glucosaminyl (N-acetyl) transferase 3, mucin type
 wingless-type MMTV integration site family member 2
 hypothetical protein FLJ23312
 KIAA1078 protein
 solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2
 hypothetical protein FLJ27274

Consensus includes gb:BF508835 /FEA=EST /DB_XREF=gi:11592133 /DB_XREF=est:U1-H-B4-acc-d-03-0-J1.s1 /CLONE=IMAGE:3085853 /UG=Hs.6217 Homo sapiens c
twisted gastrulation
deleted in azoospermia
CD3G antigen, gamma polypeptide (TIT3 complex)
Consensus includes gb:X61072.1 /DEF=Human mRNA for T cell receptor, clone IGRA17. /FEA=mRNA /PROD=T cell receptor /DB_XREF=gi:33511 /UG=Hs.272500 Huma
X61072
COP22 for noncathrin coat protein zeta-COP
Conserved gene telomeric to alpha globin cluster
dopa decarboxylase (aromatic L-amino acid decarboxylase)
cancer/testis antigen 2
CCR4-NOT transcription complex, subunit 4
KIAA1085 protein
thioredoxin reductase 1
hypothetical protein MGC10784
phorbol-like protein MDS019
Consensus includes gb:BE504197 /FEA=EST /DB_XREF=gi:9706605 /DB_XREF=est:h6611.x1 /CLONE=IMAGE:3180332 /UG=Hs.129061 Human Chromosome 16 BA
U17040
kallikrein 3, (prostate specific antigen)
5-hydroxytryptamine (serotonin) receptor 1E
hypothetical protein FLJ10842
phospholipase D1, phosphatidylcholine-specific
sprouty homolog 2 (Drosophila)
brain-specific angiogenesis inhibitor 1
alpha-2-macroglobulin
protein tyrosine phosphatase, receptor type, I polypeptide (PTPRF), interacting protein (lirin), alpha 2
paired box gene 8
caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
sarcosine dehydrogenase
EphB2
period homolog 3 (Drosophila)
Consensus includes gb:S81916.1 /DEF=phosphoglycerate kinase (alternatively spliced) human, phosphoglycerate kinase deficient patient with episodes of muscul
major histocompatibility complex, class I-like sequence
inositol 1,4,5-trisphosphate receptor, type 1
hypothetical protein FLJ20199
hypothetical protein MGC10715
phosphatidylinositol-4-phosphate 5-kinase, type I, gamma
Consensus includes gb:AL049301.1 /DEF=Homo sapiens mRNA; cDNA DKFZ564P073 (from clone DKFZ564P073) /FEA=mRNA /DB_XREF=gi:4500067 /UG=Hs.2785
neuropeptide Y receptor Y5
ras homolog gene family, member 8
hypothetical protein FLJ20551
HCGIV-6 protein
slathmin-like 3
hypothetical protein FLJ22173
Consensus includes gb:AL356954 /DEF=Human DNA sequence from clone RP11-520F24 on chromosome 13 Contains ESTs, STSs and GSSs. Contains an HNRPA1 (het
RNA binding motif protein 9
hypothetical protein FLJ11184
homolog of mouse quaking OKI (KH domain RNA binding protein)
glycoprotein M6A
Consensus includes gb:AK026879.1 /DEF=Homo sapiens cDNA: FLJ23026 fts, clone LNG01738. /FEA=mRNA /DB_XREF=gi:10439585 /UG=Hs.306863 Homo sapiens cD
hypothetical protein FLJ10950
KIAA1448 protein
SET binding factor 1
retinoic acid receptor responder (tazarotene induced) 3
ribosomal protein L37a
dopamine receptor D2
hypothetical protein
Consensus includes gb:BG485135 /FEA=EST /DB_XREF=gi:13417414 /DB_XREF=est:602503756F1 /CLONE=IMAGE:4617445 /UG=Hs.325722 Immunoglobulin kappa v
cullin 5
plexin C1
syndecin 1
gb:NM_016374.2 /DEF=Homo sapiens RBP1-like protein (BCAA), mRNA. /FEA=mRNA /GEN=BCAA /PROD=RBP1-like protein /DB_XREF=gi:13518237 /FL=gb:NM_016374
angiotensin-like 3
Consensus includes gb:AK024849.1 /DEF=Homo sapiens cDNA: FLJ21196 fts, clone COL00193. /FEA=mRNA /DB_XREF=gi:10437257 /UG=Hs.287850 Homo sapiens cD
hypothetical protein MGC3279 similar to collectins
kuronidase, alpha-L-

collagen, type VI, alpha 1
 angiotensin I converting enzyme (peptidyl-dipeptidase A) 2
 Consensus includes gb:134409.1 /DEF=Homo Sapiens (clone 83B3E13) chromosome 4p16.3 DNA fragment. /FEA=mRNA /DB_XREF=gi:600517 /UG=Hs.94799 Homo Sapiens
 butyrophilin, subfamily 1, member A1
 Consensus includes gb:AL133388 /DEF=Human DNA sequence from clone RP1-181C24 on chromosome 6p11.1-12.2. Contains the 3 end of the BMP5 gene for bone morphogenetic protein 5
 mitogen-activated protein kinase kinase 3
 a disintegrin and metalloproteinase domain 19 (meltrin beta)
 progesterone receptor membrane component 2
 thioredoxin reductase 2
 carboxyl ester lipase (bile salt-stimulated lipase)
 neurodigin 1
 gb:14455.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-reion, complete cds. /FEA=mRNA /DB_XREF=gi:398496 /FL=gb:14455.1
 TEA domain family member 4
 hypothetical protein FLJ22386
 ptp28, US snRNP 100 kd protein
 glutamate decarboxylase 1 (brain, 67kD)
 contactin associated protein-like 2
 tryptase gamma 1
 prostaglandin E receptor 3 (subtype EP3)
 RAB31, member RAS oncogene family
 dachshund homolog (Drosophila)
 hypothetical protein B
 MAP/microtubule affinity-regulating kinase like 1
 YME1-like 1 (S. cerevisiae)
 poly(A) polymerase beta (testis specific)
 zinc finger protein 207
 metalloprotein-like 5, testis-specific (testin)
 arachidonate lipooxygenase 3
 transcription termination factor, RNA polymerase I
 Consensus includes gb:AL109716.2 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 208948. /FEA=mRNA /DB_XREF=gi:9187597 /UG=Hs.200341
 dual specificity phosphatase 6
 Consensus includes gb:AF052148.1 /DEF=Homo sapiens clone 24507 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360456 /UG=Hs.169744 Homo sapiens clone 24507
 hypothetical protein FLJ11726
 hypothetical protein FLJ10488
 Consensus includes gb:BF970253 /FEA=EST /DB_XREF=gi:12337468 /DB_XREF=est:602273505F1 /CLONE=IMAGE:4361569 /UG=Hs.173684 Homo sapiens mRNA; cDNA DKFZ547E184 (from clone DKFZ547E184). /FEA=mRNA /DB_XREF=gi:9368666 /UG=Hs.8077
 zinc finger protein 200
 hypothetical protein FLJ11560
 plexin B2
 gb:NM_020887.1 /DEF=Homo sapiens hypothetical protein from clone 1659351 (LOC57397), mRNA. /FEA=mRNA /GEN=LOC57397 /PROD=hypothetical protein from clone NM_020887
 solute carrier family 22 (organic cation transporter), member 1
 myosin, heavy polypeptide 11, smooth muscle
 hypothetical protein FLJ10490
 splicing factor 3a, subunit 2, 66kD
 carbonic anhydrase XI
 Rho guanine nucleotide exchange factor (GEF) 4
 Consensus includes gb:AL049544 /DEF=Human DNA sequence from clone 24088 on chromosome 6p11.2-q12. Contains the 3 part of a gene for a novel protein similar to AL049544
 Consensus includes gb:AL390170.1 /DEF=Homo sapiens mRNA; cDNA DKFZ547E184 (from clone DKFZ547E184). /FEA=mRNA /DB_XREF=gi:9368666 /UG=Hs.8077
 melanoma antigen, family A, 3
 Consensus includes gb:BF110411 /FEA=EST /DB_XREF=gi:10940101 /DB_XREF=est:7n5202.x1 /CLONE=IMAGE:3568274 /UG=Hs.16959 ESTs
 SFRS protein kinase 2
 Cluster Incl. A1417917/ig55d04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:2112679 /clone_end=3 /gb=A1417917 /gi=4261421 /ug=Hs.20733 len=662
 regulator of G-protein signalling 5
 neuronal pentraxin 1
 potassium channel, subfamily K, member 2 (TREK-1)
 collagen, type V, alpha 1
 dynactin 4
 ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit
 KIAA1504 protein
 sialic acid binding Ig-like lectin 8
 HRI/HFB2206 protein
 ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1
 papaya protein 3
 eukaryotic translation initiation factor 1

TONDU
 glutamate rich WD repeat protein GRWD 1.14
 HYAZ2 protein 1.14
 tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains: Huntinglin Interacting protein L: transcription factor IIIA-interacting protein 1.14
 Consensus includes gb:AK000993.1 /DEF=Homo sapiens cDNA FLJ10131 fs, clone HEMBA1003041 /FEA=mRNA /DB_XREF=gi:7021986 /UG=Hs.274128 Homo sapien 1.14
 dual specificity phosphatase 10 1.14
 protein tyrosine phosphatase, non-receptor type substrate 1 1.14
 Consensus includes gb:AA757388 /FEA=EST /DB_XREF=est:2805231 /DB_XREF=IMAGE:1327098 /UG=Hs.7057 Homo sapiens C1-du-chat regl 1.14
 coagulation factor C (Limulus polyphemus homolog); coctilin 1.14
 Consensus includes gb:AA4343027 /FEA=EST /DB_XREF=gi:1995293 /DB_XREF=est:EST46865 /UG=Hs.193606 Homo sapiens PAC clone RP5-1093017 from 7q11.23-q 1.14
 Consensus includes gb:AA4343027 /FEA=EST /DB_XREF=gi:1995293 /DB_XREF=est:EST46865 /UG=Hs.193606 Homo sapiens PAC clone RP5-1093017 from 7q11.23-q 1.14
 protocadherin 22 1.14
 Consensus includes gb:AU148161 /FEA=EST /DB_XREF=gi:1009682 /DB_XREF=est:AU148161 /CLONE=MAMMA1002758 /UG=Hs.296720 Homo sapiens cDNA FLJ12 AU148161 1.14
 Interleukin 1 receptor, type I 1.14
 KIAA0328 protein 1.14
 NS1-binding protein 1.14
 KIAA0971 protein 1.14
 folate receptor 1 (adult) 1.14
 Consensus includes gb:AL139318 /DEF=Human DNA sequence from clone RP11-124B17 on chromosome 13. Contains the DCT gene for dopachrome tautomerase (dopa 1.14
 hypothetical protein 1.14
 ubiquitinatin factor E4A (UFD2 homolog, yeast) 1.14
 human homolog of Drosophila Scribble 1.14
 phospholipid acid phosphatase type 2B 1.14
 hypothetical protein FLJ14251 1.14
 Consensus includes gb:U08902.1 /DEF=Human clone 23612 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1913880 /UG=Hs.82141 Human clone 23612 mRNA sequence 1.14
 UOP-GalactosylAc beta 1,4-galactosyltransferase, polypeptide 1 1.14
 KIAA1065 protein 1.14
 leucine-rich repeat protein, neuronal 1 1.14
 a disintegrin and metalloproteinase domain 5 1.14
 Consensus includes gb:AA338945 /FEA=EST /DB_XREF=gi:1991203 /DB_XREF=est:EST43994 /UG=Hs.318825 Homo sapiens, clone IMAGE:3616574, mRNA, partial cd 1.14
 angiodotin 1 1.14
 hypothetical protein 1.14
 DNFP-566F-1524 protein 1.14
 Consensus includes gb:AK025152.1 /DEF=Homo sapiens cDNA: FLJ21499 fs, clone COL05634. /FEA=mRNA /DB_XREF=gi:10437612 /UG=Hs.306773 Homo sapiens c 1.14
 glutamate receptor, ionotropic, N-methyl D-aspartate 2B 1.14
 nucleosome assembly protein 1-like 3 1.14
 hypothetical protein FLJ13544 1.14
 Consensus includes gb:AK025363.1 /DEF=Homo sapiens cDNA: FLJ21710 fs, clone COL10087. /FEA=mRNA /DB_XREF=gi:10437865 /UG=Hs.306807 Homo sapiens c 1.14
 mesh homeo box homolog, 1 (Drosophila) 1.14
 hypothetical protein FLJ11016 1.14
 Consensus includes gb:AA1768909 /FEA=EST /DB_XREF=gi:2820147 /DB_XREF=est:nz82h01.s1 /CLONE=IMAGE:1302001 /UG=Hs.293205 ESTs. Weakly similar to BC3 1.14
 splicing factor, arginine/serine-rich 5 1.14
 hypothetical protein FLJ10496 1.14
 mannan-binding lectin serine protease 2 1.14
 HGCG.1.1 protein 1.14
 hairy/enhancer-of-split related with YRPW motif 2 1.14
 N-acetylglucosamine amidohydrolase (acid ceramidase) 1.14
 core1 UDP-galactose 4-epimerase 1.3-galactosyltransferase 1.14
 Consensus includes gb:BF987657 /FEA=EST /DB_XREF=gi:12334872 /DB_XREF=est:3597895 /DB_XREF=est:34d02.x1 /CLONE=IMAGE:1711491 /UG=Hs.89463 potassium large conductance c 1.14
 5-hydroxytryptamine (serotonin) receptor 1F 1.14
 hypothetical protein FLJ11275 1.14
 matrix metalloproteinase 9 (gelatinase B, 92kD type IV collagenase) 1.14
 centromere protein A (17kD) 1.14
 Consensus includes gb:AW974915 /FEA=EST /DB_XREF=gi:8166118 /DB_XREF=est:EST387020 /UG=Hs.116550 ESTs 1.14
 KIAA0237 gene product 1.14
 Cluster Ind. N31716:yy15c12.a1 Homo sapiens cDNA, 3 end /clone=IMAGE:271318 /clone_end=3 /gb=N31716 /gi=1152115 /ug=Hs.31754 /flen=624 1.14
 SH3-domain binding protein 5 (BTK-associated) 1.14
 hyperpolarization activated cyclic nucleotide-gated potassium channel 4 1.14
 hypothetical protein FLJ22557 1.14
 clone 1900 unknown protein 1.14
 cysteine dioxygenase, type I 1.14
 hypothetical protein FLJ20300 1.14

aryl hydrocarbon receptor nuclear translocator-like
KIAA0790 protein
Consensus includes gb:AU155621 /FEA=EST /DB_XREF=gi:11017142 /DB_XREF=PLACE1000444 /UG=Hs.306418 Homo sapiens cDNA FLJ109 AU155621
hypotheical protein FLJ10120
caudal type homeo box transcription factor 4
hypotheical protein FLJ10726
hypotheical protein
Consensus includes gb:AL137428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N1323 (from clone DKFZp761N1323) /FEA=mRNA /DB_XREF=gi:5807885 /UG=Hs.30 AL137428
hypotheical gene FLJ13072
unc-51-like kinase 1 (C. elegans)
interleukin 11
eukaryotic translation initiation factor 1A
lymphoid enhancer-binding factor 1
fibronectin 1
Consensus includes gb:AL049997.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564L102 (from clone DKFZp564L102) /FEA=mRNA /DB_XREF=gi:4884250 /UG=Hs.3083C AL278395
tumor necrosis factor receptor superfamily, member 6b, decoy
brain cell membrane protein 1
tubby super-family protein
actin related protein 2/3 complex, subunit 4 (20 kD)
similar to ubiquitin binding protein
sperm associated antigen 11
cardioembryonic antigen-related cell adhesion molecule 3
Consensus includes gb:AF16705.1 /DEF=Homo sapiens mRNA; complete cds. /FEA=mRNA /PROD=PRO2489 /DB_XREF=gi:7959908 /FL=gb:AF16705.1
dica, large (Drosophila) homolog-associated protein 2
S164 protein
defensin, alpha 5, Paneth cell-specific
glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)
CDC14 cell division cycle 14 homolog A (S. cerevisiae)
transmembrane 4 superfamily member 1
translocase of inner mitochondrial membrane 17 homolog B (yeast)
Charot-Leyden crystal protein
splicing factor 3b, subunit 3, 130kD
G protein-coupled receptor 51
gb:AF116705.1 /DEF=Homo sapiens PRO2489 mRNA, complete cds. /FEA=mRNA /PROD=PRO2489 /DB_XREF=gi:7959908 /FL=gb:AF116705.1
ubiquitin-conjugating enzyme E2 variant 1
hypotheical protein MGC2821
Consensus includes gb:U79304.1 /DEF=Human clone 23909 mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1710291 /UG=Hs.281043 Human clone 239 U79304
X-ray repair complementing defective repair in Chinese hamster cells 2
matrix metalloproteinase 11 (stromelysin 3)
prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
DiGeorge syndrome critical region gene DGS1; likely ortholog of mouse expressed sequence 2 embryonic lethal
golgi SNAP receptor complex member 2
Consensus includes gb:AK021690.1 /DEF=Homo sapiens cDNA FLJ11628 fts, clone HEMBA1004238. /FEA=mRNA /DB_XREF=gi:10432926 /UG=Hs.306610 Homo sapiens
killer cell lectin-like receptor subfamily F, member 1
zinc transporter
DKFZP434D148 protein
Rhesus blood group, CcEe antigens
Mels1, myeloid ecotropic viral integration site 1 homolog (mouse)
hypotheical protein FLJ22389
hypotheical protein FLJ20433
zinc finger protein 136 (clone pHZ-20)
Consensus includes gb:AL390143.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762C115 (from clone DKFZp762C115) /FEA=mRNA /DB_XREF=gi:5368878 /UG=Hs.2843 AL390145
BCM-like membrane protein precursor
schaefer-scute complex-like 1 (Drosophila)
Consensus includes gb:U00956.1 /DEF=Human clone KDB5.1 (CACAC)(GTG)n repeat-containing mRNA. /FEA=mRNA /DB_XREF=gi:405058 /UG=Hs.104112 Human clone
synuclein, beta
putative selenocysteine lyase
hypotheical protein FLJ20211
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1
lecithin-cholesterol acyltransferase
promyelocytic leukemia
hypotheical protein FLJ10671
Consensus includes gb:AL109730.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIIMAGE 68600. /FEA=mRNA /DB_XREF=gi:5689835 /UG=Hs.306331 AL109730

v-fos FBJ murine osteosarcoma viral oncogene homolog	BC004490	1,14
hypothetical protein ES100098	A885170	1,14
caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)	U13022	1,14
Consensus includes gb:AL137673.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434H0872 (from clone DKFZp434H0872); /FEA=mRNA /DB_XREF=gi:680784.1 /UG=Hs.30 AL137673	NM_001549	1,14
Interferon-induced protein with tetrapeptide repeats 4	NM_024903	1,14
hypothetical protein FLJ14287	NM_015456	1,14
DKFZP56850519 protein	AK024724	1,14
lysophospholipase II	NM_001740	1,14
calbindin 2, (29kD, calretinin)	NM_004833	1,14
absent in melanoma 2	NM_002847	1,14
protein tyrosine phosphatase, receptor type, N polypeptide 2	A1732802	1,14
Consensus includes gb:A1732802 /FEA=EST /DB_XREF=gi:5053915 /DB_XREF=est:ab23e08.x5 /CLONE=IMAGE:841670 /UG=Hs.292679 ESTs	NM_024845	1,14
hypothetical protein FLJ14154	M64108	1,14
collagen, type XIV, alpha 1 (undulin)	NM_014162	1,14
HSPC072 protein	NM_006754	1,14
synaptophysin-like protein	NM_001914	1,14
cytochrome b-5	NM_021944	1,14
hypothetical protein FLJ12154	NM_004445	1,14
EphB6	NM_021191	1,14
neurogenic differentiation 4	AA195124	1,14
KIAA1609 protein	NM_003528	1,14
H2B histone family, member Q	NM_003076	1,14
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 1	N73970	1,14
DKFZP5682420 protein	L26494	1,14
POU domain, class 3, transcription factor 1	D21851	1,14
leucyl-RNA synthetase, mitochondrial	AB014574	1,14
KIAA0674 protein	NM_001990	1,14
eyes absent homolog 3 (Drosophila)	AL049983	1,14
Consensus includes gb:AL049983.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D042 (from clone DKFZp564D042); /FEA=mRNA /DB_XREF=gi:4884234 /UG=Hs.2401	AV753028	1,14
transducin (beta)-like 1	NM_017826	1,14
hypothetical protein FLJ20449	NM_022894	1,14
poly(A) polymerase gamma	A1264312	1,14
truncaldehyde repeat containing 1	A1376724	1,14
COX11 homolog, cytochrome c oxidase assembly protein (yeast)	M34428	1,14
pvi-1 (murine) oncogene homolog, MYC activator	NM_004536	1,14
baculoviral IAP repeat-containing 1	AB002344	1,14
KIAA0346 protein	AL354915	1,14
Consensus includes gb:AL354915 /DEF=Human DNA sequence from clone RP11-392A19 on chromosome 13. Contains ESTs, STSs and GSSs. Contains a novel pseudoge	NM_014571	1,14
hairy/enhancer-of-split related with YRPW motif-like	AF201292	1,14
KIAA0669 gene product	AL022101	1,14
hypothetical protein similar to preferentially expressed antigen of melanoma	NM_001239	1,14
cyclin H	BC005147	1,14
FK506 binding protein 1A (12kD)	AF203728	1,14
Consensus includes gb:AF203728.1 /DEF=Homo sapiens bone morphogenetic protein BMPY (BMPY) mRNA, partial cds. /FEA=mRNA /GEN=BMPY /PROD=bone morpho	AW450751	1,14
KIAA1058 protein	BE963238	1,14
ATP-dependent RNA helicase	NM_016376	1,14
ANKHZN protein	NM_023935	1,14
hypothetical protein MGC2592	NM_000392	1,14
ATP-binding cassette, sub-family C (CFTR/MRP), member 2	AA243774	1,14
Consensus includes gb:U82303.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938329 /UG=Hs.123080 Homo	NM_017564	1,14
matrix metalloproteinase 24 (membrane-inserted)	NM_016593	1,14
hypothetical protein DKFZp434E0321	NM_004688	1,14
osylsterol 7 alpha-hydroxylase	NM_021184	1,14
N-myc (and STAT) interactor	NM_018558	1,14
Consensus includes gb:AF189251.1 /DEF=Homo sapiens cytomegalovirus partial fusion receptor mRNA, partial cds. /FEA=mRNA /PROD=cytomegalovirus partial fusion r	NM_021972	1,14
G4 protein	X63759	1,14
gamma-aminobutyric acid (GABA) receptor, theta	NM_024679	1,14
aphingosine kinase 1	AE000659	1,14
transition protein 2 (during histone to protamine replacement)	NM_014679	1,14
hypothetical protein FLJ11939	NM_014331	1,14
T cell receptor alpha locus		
KIAA0092 gene product		
solute carrier family 7, (cationic amino acid transporter, y+ system) member 11		

Consensus includes gb:A8935123 /FEA=EST/DB_XREF=gi:5873993 /DB_XREF=est:wp13n09.x1 /CLONE=IMAGE:2464789 /UG=Hs.5734.8 ESTs
CD89 antigen (p50, early T-cell activation antigen)
hypothetical protein FLJ12888
cyclin D2
ZW10 Interactor
POU domain, class 5, transcription factor 1
Tax1 (human T-cell leukemia virus type I) binding protein 1
tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)
gamma tubulin ring complex protein (76p gene)
aminopeptidase puromycin sensitive
kinonate 2-sulfatase (Hunter syndrome)
Consensus includes gb:AK024231.1 /DEF=Homo sapiens cDNA FLJ14169 fts, clone NT2RP2002056 /FEA=mRNA /DB_XREF=gi:10435558 /UG=Hs.306670 Homo sapiens
ring finger protein 1
karyopherin alpha 3 (importin alpha 4)
REV1-like (yeast)
plasmolipin
KIAA0614 protein
myosin, heavy polypeptide 4, skeletal muscle
interleukin 2 receptor, alpha
Consensus includes gb:A879383 /FEA=EST /DB_XREF=gi:5553432 /DB_XREF=est:au50b05.x1 /CLONE=IMAGE:2518161 /UG=Hs.190885 ESTs, Moderately similar to A A879383
promyelocytic leukemia
proteoglycan 1, secretory granule
Consensus includes gb:AK022686.1 /DEF=Homo sapiens cDNA FLJ12824 fts, clone NT2RM44001754 /FEA=mRNA /DB_XREF=gi:10434225 /UG=Hs.306639 Homo sapiens
chromosome 1 open reading frame 27
kidney- and liver-specific gene
PRO1598 protein
translation factor su11 homolog
collagen, type IV, alpha 6
hyaluronan binding protein 2
angiotensin-like 2
chemokine (C-C motif) receptor 9
KIAA0076 gene product
putative translation initiation factor
seven in absentia homolog 1 (Drosophila)
hypothetical protein
Kv channel interacting protein 2
hypothetical protein FLJ20331
trophinin
RAS p21 protein activator (GTPase activating protein) 3 (Ins(1,3,4,5)P4-binding protein)
erlasein homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)
apoptosis-associated tyrosine kinase
add cluster protein 33
amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)
dual specificity phosphatase 1
hypothetical protein FLJ11795
KIAA0354 gene product
Consensus includes gb:BG231551 /FEA=EST /DB_XREF=gi:12726658 /DB_XREF=est:na129h05.x1 /CLONE=IMAGE:4142457 /UG=Hs.74861 activated RNA polymerase I
KIAA0763 gene product
double C2-like domains, beta
lutealizing hormone/choriogonadotropin receptor
bromodomain-containing 1
pyrimidinergic receptor P2Y, G-protein coupled, 6
keratin 13
EHM2 gene
hypothetical protein FLJ21511
adenosine deaminase
oncomodulin
hypothetical protein FLJ23233
protein tyrosine phosphatase, non-receptor type 21
ATP-binding cassette, sub-family C (CFTR/MRP), member 5
beta-amyloid binding protein precursor
hypothetical protein FLJ20753
myoneurin

A8935123 1.14
L07555 1.14
NM_024945 1.14
NM_001759 1.14
NM_007057 1.14
NM_002701 1.14
U25801 1.14
AF021834 1.14
BF592982 1.14
AW340850 1.14
BC006170 1.14
AK024231 1.14
NM_002931 1.14
AL120704 1.14
NM_016316 1.14
AL020989 1.14
BC006270 1.14
NM_017533 1.14
NM_000417 1.14
NM_000080 1.14
J03223 1.14
AK022686 1.13
NM_017847 1.13
AB013094 1.13
NM_031270 1.13
NM_005875 1.13
AL031177 1.13
NM_004132 1.13
AF007150 1.13
AF145439 1.13
D38548 1.13
AL523310 1.13
NM_003031 1.13
NM_015371 1.13
NM_014591 1.13
NM_017788 1.13
AF349720 1.13
NM_007368 1.13
N25546 1.13
NM_004920 1.13
NM_016630 1.13
X06989 1.13
NM_004417 1.13
NM_024669 1.13
NM_014872 1.13
BG231551 1.13
NM_014869 1.13
NM_003585 1.13
NM_000233 1.13
NM_014577 1.13
NM_004154 1.13
NM_002274 1.13
NM_019114 1.13
NM_025087 1.13
Z57053 1.13
NM_006188 1.13
NM_024691 1.13
X79510 1.13
AF146074 1.13
AA012917 1.13
NM_017950 1.13
NM_016657 1.13

Consensus includes gb:AJ278883 /DEF=Homo sapiens alpha-NAC gene for nascent polypeptide-associated complex component /FEA=CDS /DB_XREF=gi:9968289 /UG= AJ278883 1.13
bone morphogenetic protein receptor, type IA NM_004329 1.13
kuronate 2-sulfatase (Hunter syndrome) AF179281 1.13
cyclin-dependent kinase 2 M68520 1.13
protein kinase, lysine deficient 1 A1788512 1.13
KIAA0602 protein AL583340 1.13
astrotactin AL031290 1.13
hydrolytic branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex, maple syrup urine disease) J03208 1.13
UDP-Gal4betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 4 AB028730 1.13
soluble carrier family 12 (sodium/chloride transporters), member 3 NM_000339 1.13
SRY (sex determining region Y)-box 10 A1367319 1.13
placenta-specific 1 NM_021796 1.13
foli-like receptor 5 AF051151 1.13
HUST checkpoint homolog (S. pombe) A1868626 1.13
BAI1-associated protein 2 NM_017450 1.13
Consensus includes gb:N92708 /FEA=EST /DB_XREF=gi:1265017 /DB_XREF=est:z644c04.s1 /CLONE=IMAGE:306438 /UG=Hs.32374 ESTs, Weakly similar to KIAA152t N92708 1.13
homoio box A6 NM_024014 1.13
NCX protein NM_016170 1.13
heterogeneous nuclear ribonucleoprotein A3 AA527502 1.13
carboxypeptidase A3 (mast cell) NM_001870 1.13
chloride channel, calcium activated, family member 2 AF043977 1.13
Consensus includes gb:AV759552 /FEA=EST /DB_XREF=gi:10917400 /DB_XREF=est:AV759552 /CLONE=MDSBCH06 /UG=Hs.226138 Homo sapiens mRNA; cDNA DK1 AV759552 1.13
diacylglycerol kinase, zeta (104kD) NM_003846 1.13
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog) Z81010 1.13
nucleic acid diphosphate linked moiety X-type motif 4 AF191654 1.13
PBX1-related 1 homeobox 2 DB1792 1.13
gb:NM_017695.1 /DEF=Homo sapiens hypothetical protein FLJ20163 (FLJ20163), mRNA, /FEA=mRNA /GEN=FLJ20163 /PRODHypothetical protein FLJ20163 /DB_XREF=NM_017695 1.13
Consensus includes gb:AK025239.1 /DEF=Homo sapiens cDNA: FLJ21586 fls, clone COL06920, /FEA=mRNA /DB_XREF=gi:10437709 /UG=Hs.321142 Homo sapiens cDNA AK025239 1.13
hypothetical protein FLJ11457 NM_024753 1.13
lectin, mannose-binding, 1 NM_005570 1.13
peptidylarginine deiminase type III NM_016233 1.13
alpha-1,4-N-acetylglucosaminyltransferase NM_016161 1.13
myosin binding protein C, fast type NM_004533 1.13
procathepsin gamma subfamily A, 3 AF152509 1.13
protein kinase Njmu-A1 NM_022344 1.13
Consensus includes gb:BE669609 /FEA=EST /DB_XREF=gi:10030150 /DB_XREF=est:7a15h11.x1 /CLONE=IMAGE:3282597 /UG=Hs.251411 Homo sapiens mRNA; cDNA BE669609 1.13
Consensus includes gb:AK024815.1 /DEF=Homo sapiens cDNA: FLJ20962 fls, clone ADSH00804, /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens cDNA AK024815 1.13
Consensus includes gb:U79277.1 /DEF=Human done 23548 mRNA sequence, /FEA=mRNA /DB_XREF=gi:1710245 /UG=Hs.71846 Human done 23548 mRNA sequence U79277 1.13
small nuclear RNA activating complex, polypeptide 4, 190kD AK023513 1.13
desmuslin AK026420 1.13
hypothetical protein FLJ12903 NM_022753 1.13
I-box and leucine-rich repeat protein 2 AF176518 1.13
phosphatidylinositol-4-phosphate 5-kinase, type II, beta NM_003559 1.13
Consensus includes gb:AI826075 /FEA=EST /DB_XREF=gi:5446746 /DB_XREF=est:wk28c04.x1 /CLONE=IMAGE:2413538 /UG=Hs.297681 serine (or cysteine) proteinase AI826075 1.13
ATPase, Cu++ transporting, beta polypeptide (Wilson disease) Consensus includes gb:A075181 /FEA=EST /DB_XREF=gi:3401772 /DB_XREF=est:oy96b04.x1 /CLONE=IMAGE:1673647 /UG=Hs.304000 ESTs, Highly similar to inositol A075181 1.13
paired box gene 8 (aniridia, keratitis) NM_000053 1.13
S100 calcium binding protein A14 (calgizarin) NM_021039 1.13
interleukin 17B NM_014443 1.13
peptidylprolyl isomerase (cyclophilin)-like 2 BC000022 1.13
homoio box B1 S49765 1.13
D component of complement (edipain) NM_001928 1.13
amyloid beta (A4) precursor-like protein 1 U48437 1.13
oxidative 3 alpha hydroxysteroid dehydrogenase; retinol dehydrogenase; 3-hydroxysteroid epimerase U89281 1.13
prostaglandin I2 (prostaglandin) synthase D38145 1.13
Consensus includes gb:AU148808 /FEA=EST /DB_XREF=gi:11010328 /DB_XREF=est:AU148808 /CLONE=NT2RM4000979 /UG=Hs.96927 Homo sapiens cDNA FLJ125t AU148808 1.13
KIAA0847 protein AW294688 1.13
Consensus includes gb:A1820796 /FEA=EST /DB_XREF=gi:5439875 /DB_XREF=est:an35h08.x5 /CLONE=IMAGE:1700703 /UG=Hs.310806 ESTs, Moderately similar to A A1820796 1.13
thymidylate synthetase BG281679 1.13
deleted in azoospermia AF271088 1.13
Consensus includes gb:AL024493 /DEF=Human DNA sequence from clone 452M16 on chromosome Xq21.1-21.33 Contains capping protein alpha subunit isoform 1 pseud AL024493 1.13
semaphorin sem2 NM_020163 1.13
Cluster Inc. AA225490.nc25501.r1 Homo sapiens cDNA /clone=IMAGE:1009129 /gb=AA225490 /gi=1846817 /ug=Hs.80115 /len=539 AA225490 1.13

goign-like protein	NM_018652	1.13
apurnic/pyrimidinic endonuclease(APEX nuclease)-like 2 protein	NM_014481	1.13
hypothetical protein PRO2730	AL137854	1.13
major histocompatibility complex, class II, DQ beta 1	M17955	1.13
DKFZP568N0319 protein	NM_015437	1.13
regulator of G-protein signaling 13	AF030107	1.13
hypothetical protein FLJ11598	NM_024966	1.13
KIAA0451 gene product	NM_014828	1.13
Consensus includes gb:U80773.1 /DEF=Human EST clone 42944 mariner transposon Hamar1 sequence. /FEA=mRNA /DB_XREF=gi:2231376 /UG=Hs.121580 Human ES U80773	NM_000194	1.13
hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome)	BE93245	1.13
I-box and WD-40 domain protein 1B	BC003617	1.13
RAS6A, member RAS oncogene family	NM_006857	1.13
formintransferase cytochrome	NM_014442	1.13
sialic acid binding Ig-like lectin 8	U92226	1.13
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	NM_006322	1.13
spindle pole body protein	NM_021622	1.13
pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 1	U42349	1.13
Pu1-like protein	U82532	1.13
Rho GDP dissociation inhibitor (GDI) gamma	BC003653	1.13
cytochrome-like nuclear factor n-pac	BE971373	1.13
cytochrome P450, subfamily I1C (mephenytoin 4-hydroxylase), polypeptide 9	AE59611	1.13
KIAA0653 protein, 87-like protein	NM_003305	1.13
transient receptor potential channel 3	NM_003305	1.13
Consensus includes gb:BF574664 /FEA=EST /DB_XREF=gi:11648376 /DB_XREF=est:602132139F1 /CLONE=IMAGE:4271792 /UG=Hs.274531 Homo sapiens mRNA; cDNA BF574664	U58111	1.13
vascular endothelial growth factor C	NM_004313	1.13
Consensus includes gb:AK025325.1 /DEF=Homo sapiens cDNA: FLJ21672 fa, clone COL09025. /FEA=mRNA /DB_XREF=gi:10437818 /UG=Hs.306801 Homo sapiens cDNA: AK025325	NM_004909	1.13
arrestin, beta 2	NM_004909	1.13
tand resistance associated gene 3	NM_004909	1.13
Consensus includes gb:AA583336 /FEA=EST /DB_XREF=gi:2367945 /DB_XREF=est:tm45108.s1 /CLONE=IMAGE:1086855 /UG=Hs.27371 Homo sapiens mRNA; cDNA: AA583336	NM_021942	1.13
hypothetical protein FLJ12716	NM_004504	1.13
HIV-1 Rev binding protein	AL041139	1.13
jerky homolog (mouse)	NM_002347	1.13
lymphocyte antigen 8 complex, locus H	NM_002347	1.13
peroxisome proliferative activated receptor, delta	BC002715	1.13
pregnancy specific beta-1-glycoprotein 6	NM_002782	1.13
phosphatidylinositol glycan, class C, pseudogene 1	AW35097	1.13
SON DNA binding protein	AA694291	1.13
huntingtin interacting protein-1-related	AB013384	1.13
nuclear receptor subfamily 2, group E, member 1	AF220532	1.13
NM23-H8	NM_016616	1.13
calcium/calmodulin-dependent protein kinase 2, beta	AF140507	1.13
hypothetical protein DKFZP434P144	AL136868	1.13
Consensus includes gb:AV704017 /FEA=EST /DB_XREF=gi:10721338 /DB_XREF=est:AV704017 /CLONE=ADBAOG04 /UG=Hs.15933 ESTs, Moderately similar to ALU7.	AV704017	1.13
solute carrier family 18 (monocarboxylic acid transporters), member 6	NM_004694	1.13
wingless-type MMTV integration site family, member 6	AY009401	1.13
Kruppel-like factor 6	NM_007250	1.13
hypothetical protein FLJ14288	NM_024764	1.13
UDP-glucuronic acid/UDP-N-acetylglucosamine dual transporter	AB044343	1.13
homeo box A7	NM_006898	1.13
phosphatidylinositol 4-kinase type II	BC003167	1.13
Consensus includes gb:U63332.1 /DEF=Human super cysteine rich protein mRNA, partial cds. /FEA=mRNA /PROD=super cysteine rich protein /DB_XREF=gi:1480862 /UC	NM_002061	1.13
glutamate-cysteine ligase, modifier subunit	BF11821	1.13
DKFZP564A122 protein	NM_014156	1.13
DKFZP564O0463 protein	NM_003951	1.13
solute carrier family 25 (mitochondrial carrier, brain), member 14	NM_003813	1.13
cartilage intermediate layer protein, nucleotide pyrophosphatase	NM_000558	1.13
hemoglobin, alpha 1	BC005325	1.13
synovial sarcoma, X breakpoint 4	NM_001825	1.13
creatine kinase, mitochondrial 2 (sarcomeric)	NM_003807	1.13
tumor necrosis factor (ligand) superfamily, member 14	BF433902	1.13
tumor necrosis factor receptor superfamily, member 11b (osteoprotegerin)	AF082537	1.13
culin 3	NM_022569	1.13
N-deacetylaseN-sulfotransferase 4		

NAG-7 protein
 keratocan
 hypothetical protein PRO2133
 mitogen-activated protein kinase 13
 hypothetical protein FLJ11608
 v-maf musculoaponeurotic fibrosarcoma oncogene homolog F (avian)
 matrix metalloproteinase 3 (stromelysin 1, progelatinase)
 Consensus includes gb:AL137270.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434G0814 (from clone DKFZp434G0814). /FEA=mRNA /DB_XREF=gj:6807705 /UG=Hs.39 AL137270
 Consensus includes gb:AL137270.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434G0814 (from clone DKFZp434G0814). /FEA=mRNA /DB_XREF=gj:6807705 /UG=Hs.39 AL137270
 Consensus includes gb:N1149 /FEA=EST /DB_XREF=gj:1444476 /UG=Hs.279639 Homo sapiens mRNA; cDNA DK N81149
 carboxylate (chondroin & keratan) sulfotransferase 3
 serine/threonine kinase 23
 Consensus includes gb:AI186739 /FEA=EST /DB_XREF=gj:3737377 /DB_XREF=est:ae79601.x1 /CLONE=IMAGE:1745184 /UG=Hs.326416 Homo sapiens mRNA; cDNA AI186739
 axonal transport of synaptic vesicles
 caldinin/calactinin-related polypeptide, alpha
 hypothetical protein FLJ11271
 pumilio (Drosophila) homolog 2
 topoisomerase (DNA) II beta (180kD)
 ATP-binding cassette, sub-family F (GCM20), member 2
 hypothetical protein
 KIA00737 gene product
 potassium voltage-gated channel, subfamily F, member 1
 junction plakoglobin
 Consensus includes gb:AU144309 /FEA=EST /DB_XREF=gj:11005830 /DB_XREF=est:AU144309 /CLONE=HEMBA1001533 /UG=Hs.55830 Homo sapiens cDNA FLJ1001
 KIA0217 protein
 hypothetical protein MGC10520
 CGL-53 protein
 high-mobility group (nonhistone chromosomal) protein 17-like 1
 sorting nexin 13
 cAMP responsive element binding protein-like 2
 Consensus includes gb:AI417917 /FEA=EST /DB_XREF=gj:4261421 /DB_XREF=est:ig55604.x1 /CLONE=IMAGE:2112679 /UG=Hs.20733 Homo sapiens cDNA: FLJ22356 /AI417917
 hypothetical protein FLJ14106
 myelin basic protein
 Koppel-associated box protein
 Consensus includes gb:BF448596 /FEA=EST /DB_XREF=gj:11514774 /DB_XREF=est:7n90504.x1 /CLONE=IMAGE:3571902 /UG=Hs.290484 ESTs
 exportin, tRNA (nuclear export receptor for tRNAs)
 ubiquitin specific protease 2
 myosin regulatory light chain
 hypothetical protein FLJ46203.2
 KIA0280 protein
 microtubule-associated protein tau
 Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fls, clone COL09829. /FEA=mRNA /DB_XREF=gj:10437851 /UG=Hs.306804 Homo sapiens cDNA: FLJ22356 /AI417917
 neuropilin 2
 chorionic somatomammotropin hormone 2
 MRE11 meiotic recombination 11 homolog A (S. cerevisiae)
 KIA0255 gene product
 PTPN13-like, Y-linked
 phosphatase and tensin homolog (mutated in multiple advanced cancers 1)
 KIA0808 gene product
 peptidylprolyl isomerase D (cyclophilin D)
 Consensus includes gb:AT32885 /FEA=EST /DB_XREF=gj:5053998 /DB_XREF=est:oe64d04.x5 /CLONE=IMAGE:1416391 /UG=Hs.142907 Human BRCA2 region, mRNA /AI732885
 SecE1 alpha form 2
 regulator of G-protein signalling 9
 hypothetical protein FLJ23505
 hypothetical protein FLJ13273
 FYVE and coiled-coil domain containing 1
 hypothetical protein FLJ11710
 sarcoglycan, epsilon
 lymphocyte antigen 9
 Consensus includes gb:AC006986 /DEF=Homo sapiens BAC clone RP11-155J5 from Y /FEA=CDS /DB_XREF=gj:4753246 /UG=Hs.283908 Homo sapiens BAC clone RP1AC006986
 keratin 17
 FSHD region gene 1
 actin related protein 23 complex, subunit 2 (34 kD)
 Consensus includes gb:AK022418.1 /DEF=Homo sapiens cDNA FLJ12356 fls, clone MAMMA1002339. /FEA=mRNA /DB_XREF=gj:10433811 /UG=Hs.306635 Homo sapiens cDNA: FLJ22356 /AI732885

hypothetical protein FLJ12389 similar to octoacetyl-CoA synthetase	NM_023928	1.13
potassium channel, subfamily K, member 15 (TASK-5)	NM_022356	1.13
histidine rich calcium binding protein	NM_002152	1.13
eukaryotic translation initiation factor 4E binding protein 2	BC005057	1.13
ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase: transferase B, alpha 1-3-galactosyltransferase)	U15197	1.13
Consensus includes gb:AK024543.1 /DEF=Homo sapiens cDNA: FLJ20890 fs, clone ADKA03323. /FEA=mRNA /DB_XREF=gi:10436848 /UG=Hs.306686 Homo sapiens c AK024543	NM_014677	1.13
KIAA0751 gene product	NM_000796	1.13
dopamine receptor D3	AA181053	1.13
pervin, beta	AI189589	1.13
Consensus includes gb:AA330995 /FEA=EST /DB_XREF=gi:2273701 /DB_XREF=est:n07a02.s1 /CLONE=IMAGE:985610 /UG=Hs.23729 Homo sapiens clone 24405 mRI	NM_002250	1.13
hypothetical protein DKFZp434E1723	BC001279	1.13
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4	NM_004143	1.13
gb:BC001278.1 /DEF=Homo sapiens. Similar to KIAA0628 gene product, clone MGC:5129, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0628 gene product	NM_002299	1.13
Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 1	AI762021	1.13
platelet alpha 1 (ectodermal dysplasia/skin fragility syndrome)	NM_001756	1.13
synaptobin, beta 2 (dystrophin-associated protein A1, 59kD, basic component 2)	AC003079	1.13
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 6	NM_014717	1.13
ankyrin repeat and SOCS box-containing 4	AF070528	1.13
KIAA0390 gene product	NM_004912	1.13
mucosa associated lymphoid tissue lymphoma translocation gene 1	BE466926	1.13
cerebral cavernous malformations 1	NM_003159	1.13
serine/threonine kinase 9	NM_003069	1.13
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1	NM_016147	1.13
protein phosphatase methyltransferase-1	NM_017701	1.13
Rho GTPase activating protein 8	NM_003165	1.13
synaptin binding protein 1	AW277253	1.13
adenylyate kinase 2	R37427	1.13
Consensus includes gb:R37427 /FEA=EST /DB_XREF=gi:794883 /DB_XREF=est:y55g12.s1 /CLONE=IMAGE:25997 /UG=Hs.6524 Homo sapiens clone IMAGE 25997	X07247	1.13
alkaline phosphatase, placental-like 2	NM_000260	1.13
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	NM_025214	1.13
CTCL tumor antigen se57-1	AI829170	1.13
gokjin-67	AI539305	1.13
chromosome 21 open reading frame 2	BC002456	1.13
voltage-dependent anion channel 3	NM_001952	1.13
E2F transcription factor 6	NM_000376	1.13
vitamin D (1,25-dihydroxyvitamin D3) receptor	AA071195	1.13
chloride channel 4	BC000189	1.13
olfactomedin 1	NM_002171	1.13
interferon, alpha 10	AW029619	1.13
cytoskeleton-associated protein 4	NM_017682	1.12
hypothetical protein FLJ20132	AI041520	1.12
fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor)	NM_006700	1.12
FLN29 gene product	NM_000055	1.12
butyrylcholinesterase	AJ000008	1.12
phosphoinositide-3-kinase, class 2, gamma polypeptide	BE567032	1.12
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	BF875004	1.12
poly(A) binding protein, nuclear 1	AA747379	1.12
calcitonin-related polypeptide, beta	NM_015893	1.12
plasmidipin	AA719787	1.12
olfactory receptor, family 7, subfamily E, member 13 pseudogene	AV702994	1.12
HSPCO34 protein	AI133262	1.12
Consensus includes gb:AL133262 /DEF=Human DNA sequence from clone RP1-319022 on chromosome 6 Contains part of the gene for a CDC5-like protein, the gene for	M33987	1.12
carbonic anhydrase I	AF031469	1.12
major histocompatibility complex, class I-like sequence	W07700	1.12
hypothetical protein MGC13098	K59217	1.12
collagen, type XIII, alpha 1	NM_002120	1.12
major histocompatibility complex, class II, DO beta	NM_005590	1.12
Consensus includes gb:AL523076 /FEA=EST /DB_XREF=gi:12786569 /DB_XREF=est:AL523076 /CLONE=CS00C001Y112 (3 prime) /UG=Hs.82503 H.sapiens mRNA for	NM_002012	1.12
MRE11 meiotic recombination 11 homolog A (S. cerevisiae)	NM_004815	1.12
fragile histidine triad gene	AI734124	1.12
PTPL1-associated RhoGAP 1		
ribosomal protein S14		

immunoglobulin kappa constant
Consensus includes gb:AV700132 /FEA=EST /DB_XREF=gi:10302103 /DB_XREF=est:AV700132 /CLONE=GKCGSE03 /UG=Hs.295923 seven in absentia (Drosophila) hc:AV700132
U12543
KIAA0288 gene product
cleft lip and palate associated transmembrane protein 1
hypothetical protein
protease, serine, 7 (enterokinase)
hepsin (transmembrane protease, serine 1)
ATP-binding cassette, sub-family A (ABC1), member 2
bone morphogenetic protein receptor, type IA
Cas-Bc-M (murine) ectropic retroviral transforming sequence b
UDP-glucose ceramide glucosyltransferase-like 1
hypothetical protein DKFZp566133
signal sequence receptor, gamma (translocon-associated protein gamma)
RAS, member RAS oncogene family
damage-specific DNA binding protein 2 (48kD)
interleukin 9
hypothetical protein IMAGE 106914
Consensus includes gb:AL060215.1 /DEF=Homo sapiens mRNA; cDNA DKFZp568J0323 (from clone DKFZp568J0323) /FEA=mrna /DB_XREF=gi:5262706 /UG=Hs.102 AL060215
Cluster ind. A1830563;g51c03.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2406340 /clone_and=3 /gb=A1830563 /gi=5451234 /ug=Hs.120858 /len=556
hypothetical protein DKFZp434K1172
centaurin, alpha 1
solite carrier family 7 (cationic amino acid transporter, y+ system), member 2
KIAA0387 protein
amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3
growth hormone 1
actin, alpha 3
ATPase, Na+/K+ transporting, beta 2 polypeptide
tumor necrosis factor receptor superfamily, member 8
hypothetical protein similar to tenascin-R
hypothetical protein FLJ13213
forkhead box F2
runt-related transcription factor 3
hypothetical protein MGC10646
DNA-dependent protein kinase catalytic subunit-interacting protein 2
Consensus includes gb:AJ252550 /DEF=Homo sapiens partial CK gene for glycerol kinase, exon 1 (glycerol kinase deficiency case) /FEA=CDS_1 /DB_XREF=gi:5834425 /UG=Hs.201817 Homo sapiens cDNA FLJ121 AU146949
KIAA0228 gene product
glutamate receptor, ionotropic, delta 2
hemochromatosis
DNA (cytosine-5-)-methyltransferase 3 alpha
NAD kinase
centrin, EF-hand protein, 1
KIAA0815 gene product
Consensus includes gb:AU146949 /FEA=EST /DB_XREF=gi:11008470 /DB_XREF=est:AU146949 /CLONE=HEM8B1001952 /UG=Hs.301817 Homo sapiens cDNA FLJ121 AU146949
KIAA1056 protein
KIAA0712 gene product
solite carrier family 25 (mitochondrial carrier; Graves disease autoantigen), member 18
hydroxyprostaglandin dehydrogenase 15-(NAD)
interleukin 6 (interferon, beta 2)
cyclic AMP-regulated phosphoprotein, 21 kD
hypothetical protein FLJ20151
signal transducer and activator of transcription 1, 91kD
C-terminal binding protein 2
Kv channel-interacting protein 1
calcium channel, voltage-dependent, beta 3 subunit
UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 6
protein phosphatase 1, regulatory subunit 3D
DKFZP434L243 protein
KIAA0790 protein
Small evolutionarily conserved RNA, resembling C/D box small nuclear
Consensus includes gb:AU147317 /FEA=EST /DB_XREF=gi:11008838 /DB_XREF=est:AU147317 /CLONE=MAMMA1000340 /UG=Hs.287491 Homo sapiens cDNA FLJ12 AU147317
synovelin
hypothetical protein FLJ10628

dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A	U52373	1.12
Consensus includes gb:U79273.1 /DEF=Human clone 23933 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710239 /UG=Hs.239483 Human clone 23933 mRNA sequenc	U79273	1.12
KIAA0239 protein	A1735639	1.12
hypothetical protein	AK027245	1.12
alcohol dehydrogenase 1B (class I), beta polypeptide	M24317	1.12
LW-1	NM_016153	1.12
Kuppel-like factor 12	NM_007249	1.12
ERGL protein; ERGIC-53-like protein	NM_021819	1.12
meningioma expressed antigen 5 (hyaluronidase)	AK002091	1.12
Consensus includes gb:AV661138 /FEA=EST /DB_XREF=gi:9882152 /DB_XREF=est:AV661138 /CLONE=GLCGPA11 /UG=Hs.282712 ESTs	AV661138	1.12
hypothetical protein PR01094	NM_018575	1.12
methionine adenosyltransferase I, alpha	NM_000428	1.12
hypothetical SBB03 protein	NM_005785	1.12
proteasome gamma subfamily C, 3	BC006439	1.12
hypothetical protein FLJ20147	NM_017687	1.12
Consensus includes gb:Z69744 /DEF=H.sapiens ALL-1 gene exon 1 (and joined coding region) /FEA=CDS /DB_XREF=gi:1490270 /UG=Hs.199160 myeloidlymphoid or mlt: Z69744	AK025271	1.12
Consensus includes gb:AK025271.1 /DEF=Homo sapiens cDNA: FLJ21618 fs, clone COL07487. /FEA=mRNA /DB_XREF=gi:10437753 /UG=Hs.306790 Homo sapiens cD: AK025271	AF067286	1.12
TAP binding protein (tapasin)	AF067286	1.12
heat shock 70kD protein 8B (mortalin-2)	A4927701	1.12
hypothetical protein FLJ21120	NM_024767	1.12
Cluster incl. AF007155: Homo sapiens clone 23763 unknown mRNA, partial cds /cds=(0,314) /gb=AF007155 /gi=2852635 /ug=Hs.186894 /len=1369	AF007155	1.12
spindlin-like	AF356353	1.12
SEC24 related gene family, member D (S. cerevisiae)	AK000709	1.12
Rho GTPase activating protein 1	BG468434	1.12
ubiquitin specific protease 7 (herpes virus-associated)	A1160440	1.12
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9	NM_014358	1.12
actinin	NM_006576	1.12
p53-regulated apoptosis-inducing protein 1	NM_022112	1.12
collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant)	AI813758	1.12
cystatin D	NM_001900	1.12
replication factor C (activator 1) 3 (38kD)	BC000149	1.12
cytochrome P450, subfamily IID (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6	X07618	1.12
chromodomain protein, Y chromosome, 1	AF080597	1.12
miS homolog 4 (E. coli)	AF104243	1.12
cyclin D1 (PRAD1; parathyroid adenomatosis 1)	Z23022	1.12
collagen, type XVIII, alpha 1	AF018081	1.12
pre-TiNK cell associated protein	NM_021981	1.12
annexin A4	BC000182	1.12
fetal Alzheimer antigen	NM_004459	1.12
zinc finger protein 198	AL136621	1.12
N-deacetylaseN-sulfotransferase (heparan glucosaminyl) 3	NM_004784	1.12
RNA guanylyltransferase and 5'-phosphatase	AB012143	1.12
lysosomal apyrase-like 1	NM_004901	1.12
serine/threonine kinase 17a (apoptosis-inducing)	NM_004760	1.12
ribonuclease L (2',5'-oligoadenylate synthetase-dependent)	NM_021133	1.12
serine/threonine-protein kinase PRP4 homolog	AB011108	1.12
hypothetical protein FLJ20527	NM_017863	1.12
hypothetical protein FLJ20730	NM_017845	1.12
ELK4, ETS-domain protein (SRF accessory protein 1)	NM_001973	1.12
NICE-1 protein	NM_019060	1.12
CD97 antigen	NM_001784	1.12
hypothetical protein FLJ10902	NM_018268	1.12
Consensus includes gb:AA224446 /FEA=EST /DB_XREF=gi:1844989 /DB_XREF=est:zt16803.1 /CLONE=IMAGE:663532 /UG=Hs.311561 ESTs, Weakly similar to CYA7	AA224446	1.12
death-associated protein 6	BE965715	1.12
small nuclear RNA activating complex, polypeptide 4, 190kD	NM_003086	1.12
solute carrier family 6 (neurotransmitter transporter, GABA), member 1	A003579	1.12
hemoglobin, alpha 2	AF105974	1.12
lectin, galactoside-binding, soluble, 3 binding protein	NM_005567	1.12
cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 7	M28873	1.12
Consensus includes gb:AU150681 /FEA=EST /DB_XREF=gi:11012212 /DB_XREF=est:AU150681 /CLONE=NT2RP2003367 /UG=Hs.153321 Homo sapiens cDNA FLJ105	AU150681	1.12
regulator of G-protein signalling 12	AB24113	1.12
beta tubulin 1, class VI	NM_030773	1.12
hypothetical protein FLJ11565	NM_024657	1.12

KIAA0620 protein
Consensus includes gb:U80768.1 /DEF=Human EST clone 22453 mariner transposon Hsmar1 sequence. /FEA=mRNA /DB_XREF=gi:2231369 /UG=Hs.13252 Human EST U80766
Consensus includes gb:AF007147.1 /DEF=Homo sapiens clone 23712 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852625 /UG=Hs.113166 Homo sapiens clone 23712 AF007147
DKFZP564K247 protein
KIAA1031 protein
chondroectoderm
organic anion transporter polypeptide-related protein 4
hypothetical protein FLJ20651
serine (or cysteine) proteinase inhibitor, clade I (neuroserpin), member 1
hypothetical protein FLJ2341
arachidonate 5-lipoxygenase
proteoglycan 3
glucagon-like peptide 1 receptor
keratin 15
nidogen (enactin)
nuclear transcription factor, X-box binding 1
cyclin-dependent kinase inhibitor 1C (p57, Kip2)
Consensus includes gb:U79300.1 /DEF=Human clone 23629 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710285 /UG=Hs.135587 Human clone 23629 mRNA sequence U79300
GDNF family receptor alpha 4
solute carrier family 22 (organic anion transporter), member 6
cytochrome P450, subfamily IVB, polypeptide 1
insulin receptor
ATPase, Na⁺/K⁺ transporting, alpha 3 polypeptide
Consensus includes gb:AK023891.1 /DEF=Homo sapiens cDNA FLJ13829 fis. clone THYRO1000825. /FEA=mRNA /DB_XREF=gi:10435964 /UG=Hs.87819 Homo sapiens, clone MGC:249; A829961
DKFZP564N1363 protein
lipin 1
prolactin receptor
mesenchymal stem cell protein DSCD75
angiotensin receptor-like 1
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 5
Consensus includes gb:A856610 /FEA=EST /DB_XREF=gi:4740589 /DB_XREF=esttt46n03.x1 /CLONE=IMAGE:2243861 /UG=Hs.87819 Homo sapiens, clone MGC:249; A829961
CD7 antigen (p41)
prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
mucin 6, gastric
unc-51-like kinase 2 (C. elegans)
ras homolog gene family, member A
MHC class II transactivator
eosinophil chemotactic cytokine
calbindin 1, (28KD)
gamma-aminobutyric acid (GABA) A receptor, gamma 2
chromosome condensation 1-like
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)
BCL2-like 2
Ewing sarcoma breakpoint region 1
highly charged protein
Wolfram syndrome 1 (wolframin)
CTG28 alternate open reading frame
integrin, beta 3 (platelet glycoprotein IIb, antigen CD61)
pleckstrin homology-like domain, family A, member 1
NK-2 (Drosophila) homolog 8
hypothetical protein FLJ22054
zinc finger protein 295
hypothetical protein FLJ11535
hypothetical protein FLJ20595
H1 histone family, member 1
5-hydroxytryptamine (serotonin) receptor 4
membrane-spanning 4-domains, subfamily A, member 12 4-domains, subfamily A, member 7
myristoylated alanine-rich protein kinase C substrate
RNA-binding protein gene with multiple splicing
solute carrier family 5 (sodium/glucose cotransporter), member 1
synaptobrevin 1A (brain)
Wiskott-Aldrich syndrome-like
testis specific protein 1 (probe H4-1 p3-1)

aldolase B, fructose-bisphosphate
 staufen, RNA binding protein, homolog 2 (*Drosophila*)
 Consensus includes gb:AK024128.1 /DEF=Homo sapiens cDNA FLJ14066 fs, clone HEMBB1001197, /FEA=mrna /DB_XREF=gi:10436433 /UG=Hs.306665 Homo sapie
 plasmidogen activator, tissue
 raiA binding protein 1
 deathless, autosomal dominant 5
 Consensus includes gb:AF007143.1 /DEF=Homo sapiens clone 23738 mRNA sequence, /FEA=mrna /DB_XREF=gi:2852820 /UG=Hs.28205 Homo sapiens clone 23738
 Consensus includes gb:AF007143.1 /FEA=EST /DB_XREF=gi:5112883 /DB_XREF=est /wg0204.x1 /CLONE=IMAGE:2363935 /UG=Hs.186970 ESTs
 protein kinase C, beta 1
 Consensus includes gb:AA115963 /FEA=EST /DB_XREF=gi:1670996 /DB_XREF=est /zm78d11.s1 /CLONE=IMAGE:531765 /UG=Hs.323423 ESTs, Moderately similar to B AA115963
 G protein-coupled receptor 49
 basic transcription element binding protein 1
 vav 3 oncogene
 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 4
 KIAA0830 protein
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3
 hypothetical protein FLJ20739
 v-ros UR2 sarcoma virus oncogene homolog 1 (avian)
 histone deacetylase 3
 ribosomal protein S6 kinase, 90kD, polypeptide 3
 hypothetical protein MGC4161
 neuropilin Y receptor Y2
 aldolase B, fructose-bisphosphate
 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein)
 deleted in azoospermia 2
 poly(IC) binding protein 4
 hypothetical protein FLJ23053
 ACO for serine protease homologue
 plein A2
 hypothetical protein FLJ20275
 suppressor of Ty 3 homolog (S. cerevisiae)
 cytochrome P450 polypeptide 43
 cholinergic receptor, nicotinic, beta polypeptide 3
 KIAA0428 gene product
 H2B histone family, member N
 hypothetical protein MGC4022
 hypothetical protein FLJ13181
 defensin, alpha 1, myeloid-related sequence
 CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)
 apolipoprotein C-IV
 zinc finger protein 41
 poliovirus receptor
 COQ7 coenzyme Q, 7 homolog ubiquinone (yeast)
 polymerase (RNA) II (DNA directed) polypeptide B (140kD)
 KIAA1055 protein
 solute carrier family 25, member 13 (citrin)
 RNA binding motif protein 3
 BAI1-associated protein 1
 solute carrier family 8 (sodium/calcium exchanger), member 1
 Consensus includes gb:AK025388.1 /DEF=Homo sapiens cDNA: FLJ21735 fs, clone COLF3350, /FEA=mrna /DB_XREF=gi:10437892 /UG=Hs.306808 Homo sapiens c
 B cell RAG associated protein
 chromosome 21 open reading frame 108
 hypothetical protein FLJ23394
 Consensus includes gb:AK021636.1 /DEF=Homo sapiens cDNA FLJ11774 fs, clone HEMBA1005884, /FEA=mrna /DB_XREF=gi:10433104 /UG=Hs.293197 Homo sapie
 synixin 18
 eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD)
 antyrim repeat, family A (RFXANK-like), 2
 KIAA0804 protein
 synovial sarcoma translocation gene on chromosome 18-like 2
 KIAA0998 protein
 syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)
 dual specificity phosphatase 2
 prostaglandin 12 (prostaglycin) synthase

Consensus includes gb:AK021987.1 /DEF=Homo sapiens cDNA FLJ11925 fs, clone HEMBB1000354, /FEA=mrna /DB_XREF=gi:10433296 /UG=Hs.191158 Homo sapie AK021987
 e1aC homolog 2 (E. coli) AF304370 1.11
 zinc finger protein 79 (p17) AA284828 1.11
 insulin-like growth factor 1 (somatomedin C) M37484 1.11
 phosphatidic acid phosphatase type 2B AV725664 1.11
 hypothetical protein FLJ20287 NM_017746 1.11
 zinc finger protein 308 BC008118 1.11
 transcription factor 7-like 2 (T-cell specific, HMG-box) AJ270770 1.11
 selectin E (endothelial adhesion molecule 1) NM_000450 1.11
 huntingtin interacting protein-1-related AB013384 1.11
 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) M84445 1.11
 crystallin, gamma C NM_020989 1.11
 N-ethylmaleimide-sensitive factor attachment protein, alpha BC001165 1.11
 aldehyde dehydrogenase 3 family, member A2 BC002430 1.11
 hypothetical protein from clone 24828 AF131833 1.11
 histidyl-RNA synthetase-like U18937 1.11
 ribosomal protein L10 BC003358 1.11
 regulator of G-protein signalling 12 AW170602 1.11
 Consensus includes gb:AA733079 /FEA=EST /DB_XREF=gi:2754438 /DB_XREF=est:zg78d12 s1 /CLONE=IMAGE:399575 /UG=Hs.285553 Homo sapiens, clone IMAGE: X62048 1.11
 WEE1+ homolog (S. pombe) NM_020119 1.11
 hypothetical protein FLB6421 BG500087 1.11
 Ras-GTPase-activating protein SH3-domain-binding protein NM_012102 1.11
 arginine-glutamic acid dipeptide (RE) repeats NM_017705 1.11
 hypothetical protein FLJ20190 NM_015652 1.11
 DKFZP564P1916 protein L36874 1.11
 synuclein, alpha (non A4 component of amyloid precursor) NM_000638 1.11
 vitronectin (serum spreading factor, somatomedin B, complement S-protein) NM_006374 1.11
 Consensus includes gb:S83374.1 /DEF=glutamate transporter II variant BH8GT IIB (5 region) human, brain and spinal cord, mRNA Partial Mutant, 129 nt, /FEA=mrna /DI S83374 1.11
 LIM homeobox protein 3 NM_006054 1.11
 Consensus includes gb:AL031178 /DEF=Human DNA sequence from clone 341E18 on chromosome 6p11.2-12.3. Contains a Serine/Threonine Protein Kinase gene (presu AF156888 1.11
 RA051-like 3 (S. cerevisiae) AL031178 1.11
 Kruppel-type zinc finger protein AF034956 1.11
 Consensus includes gb:BF594294 /FEA=EST /DB_XREF=gi:11686618 /DB_XREF=est:7h9c12.x1 /CLONE=IMAGE:3322198 /UG=Hs.287335 ESTs, Highly similar to AC1 BF594294 N21541 1.11
 DKFZP586C1820 protein AA524525 1.11
 olfactory receptor, family 2, subfamily H, member 1 AJ302604 1.11
 phospholipid scramblase 3 NM_020360 1.11
 I-complex 11 (a murine top homolog) NM_018679 1.11
 ATPase, Ca++ transporting, ubiquitous AF068220 1.11
 transcription factor (SMIF gene) NM_018403 1.11
 uracil-DNA glycosylase 2 BC004877 1.11
 major histocompatibility complex, class II, DR beta 3 AA807056 1.11
 hypothetical protein NM_019038 1.11
 suppressor of white spot homolog 2 NM_007056 1.11
 KIAA0303 protein NM_007056 1.11
 succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD AB002301 1.11
 ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin) D49737 1.11
 ATPase, Ca++ transporting, ubiquitous L35594 1.11
 myosin IIIA NM_005173 1.11
 Cluster Ind. N92708:zb44cd04.s1 Homo sapiens cDNA, 3 and /clone=IMAGE:306438 /clone_end=3 /gb=N92708 /gi=1265017 /ug=Hs.32374 /len=668 NM_017433 1.11
 amelogenin (Y chromosome) N92708 1.11
 solute carrier family 3 (cystine, dibasic and neutral amino acid transporters, activator of cystine, dibasic and neutral amino acid transport), member 1 NM_001143 1.11
 transcription factor-like 4 M85548 1.11
 small nuclear ribonucleoprotein polypeptide G NM_003096 1.11
 ubiquitin-like 3 NM_007106 1.11
 MAD5 box transcription enhancer factor 2, polypeptide B (myocyte enhancer factor 2B) BC004449 1.11
 glutathione S-transferase theta 2 NM_000854 1.11
 solute carrier family 12 (potassium/chloride transporters), member 6 NM_005135 1.11
 proprotein convertase subtilisin/kexin type 5 NM_008200 1.11
 Consensus includes gb:AK024887.1 /DEF=Homo sapiens cDNA: FLJ21214 fs, clone COL00523, /FEA=mrna /DB_XREF=gi:10437277 /UG=Hs.306710 Homo sapiens cC AK024887 1.11
 Consensus includes gb:AK021474.1 /DEF=Homo sapiens cDNA FLJ11412 fs, clone HEMBA1000876, /FEA=mrna /DB_XREF=gi:10432868 /UG=Hs.289094 Homo sapie AK021474 1.11
 aminocyclase 1 NM_000666 1.11
 hypothetical protein DKFZp434F0318 NM_030817 1.11

muslin 5, subtypes A and C, tracheobronchial/gastric	AW192795	1,11
Williams-Beuren syndrome chromosome region 5	BC006080	1,11
guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1	NM_021955	1,11
solute carrier family 24 (sodium/potassium/calcium exchanger), member 3	R62432	1,11
hypothetical protein FLJ12649	NM_024597	1,11
retinoblastoma 1 (including osteosarcoma)	NM_000321	1,11
cytochrome P450, subfamily IIF, polypeptide 1	NM_000774	1,11
mesoderm specific transcript homolog (mouse)	NM_002402	1,11
distal-less homeobox 4	AF254115	1,11
mal, T-cell differentiation protein	NM_002371	1,11
hypothetical protein FLJ11715	NM_024564	1,11
C-type lectin-like receptor-2	NM_016509	1,11
securin related 6 homolog (mouse)-like	AB041736	1,11
SMC1 structural maintenance of chromosomes 1-like 1 (yeast)	D80000	1,11
immunoglobulin kappa constant	BG540628	1,11
signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	AI571896	1,11
dihydrolipoamide S-acyltransferase (E2 component of pyruvate dehydrogenase complex)	AW299740	1,11
hypothetical protein FLJ10932	NM_018277	1,11
basic transcription factor 3, like 1	NM_001208	1,11
Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mRNA /DB_XREF=gj:5915893 /U AF005487	NM_022042	1,11
solute carrier family 28 (sulfate transporter), member 1	AU118165	1,11
zinc finger protein 37a (KOX 21)	X99142	1,11
keratin, hair, basic, 6 (monilebrix)	NM_005814	1,11
glycoprotein A33 (transmembrane)	AL049938	1,11
DKF ZP564P1916 protein	BC005369	1,11
egl nine homolog 1 (C. elegans)	NM_014945	1,11
KIAA0843 protein	U43143	1,11
frs-related tyrosine kinase 4	NM_017555	1,11
egl nine homolog 2 (C. elegans)	AI978718	1,11
KIAA0365 gene product	NM_001921	1,11
dCMP deaminase	NM_000102	1,11
cytochrome P450, subfamily XVII (steroid 17-alpha-hydroxylase), adrenal hyperplasia	BG230586	1,11
solute carrier family 7 (cationic amino acid transporter, y+ system), member 6	NM_018274	1,11
lensin	NM_002148	1,11
homeo box D10	AA209420	1,11
zinc finger protein 75 (D8C6)	AB006828	1,11
KIAA0280 protein	NM_003481	1,11
ubiquitin specific protease 5 (isopeptidase T)	AL042588	1,11
paternally expressed 3	AI433712	1,11
methylnalonyl Coenzyme A mutase	NM_005372	1,11
v-mos Moloney murine sarcoma viral oncogene homolog	NM_022036	1,11
G protein-coupled receptor, family C, group 5, member C	AU146004	1,11
KIAA0231 protein	AW139369	1,11
zinc finger protein 291	AW295066	1,11
Consensus includes gb:AW295066 /FEA=EST /DB_XREF=gj:6701702 /DB_XREF=est:UI-HBW0-alt-b-09-0-UI.s1 /CLONE=IMAGE:2730209 /UG=Hs.255387 ESTs	NM_003514	1,11
H2A histone family, member N	AB046836	1,11
KIAA1816 protein	NM_021959	1,11
protein phosphatase 1, regulatory (inhibitor) subunit 11	NM_024774	1,11
hypothetical protein FLJ21924	NM_012134	1,11
telomodin 1 (smooth muscle)	NM_012134	1,11
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myelin transcription factor 1-like	NM_005291	1,11
G protein-coupled receptor 17	AI189609	1,11
RAS2, member RAS oncogene family	AI189609	1,11
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albumin	AK026411	1,11
aldolase B, fructose-bisphosphate	NM_019040	1,11
hypothetical protein	NM_019040	1,11
Consensus includes gb:AK000793.1 /DEF=Homo sapiens cDNA FLJ20286 fs, clone HEP04358. /FEA=mRNA /DB_XREF=gj:7020278 /UG=Hs.306391 Homo sapiens cDN	AK000293	1,11
hypothetical protein FLJ10357	R42449	1,11
gb:Z25434.1 /DEF=H sapiens protein-serine/threonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gj:405746 /FL=gb.Z25	Z25434	1,11
stromal cell-derived factor 1	NM_000609	1,11
UBX domain-containing 2	AI927512	1,11

glucagon-like peptide 1 receptor	U01157	1,11
H4 histone family, member K	NM_003546	1,11
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 2	AF162274	1,11
CSE1 chromosome segregation 1-like (yeast)	AF053640	1,11
ectodysplasin 1, anhidrotic receptor	NM_022336	1,11
granulysin 1, anhidrotic receptor	NM_012198	1,11
grancalcin, EF-hand calcium binding protein	NM_018268	1,11
hypothetical protein FLJ10904	NM_006841	1,11
transporter protein, system N1 Na+ and H+-coupled glutamine transporter	AA625133	1,11
hypothetical protein FLJ2439	BC002809	1,11
down-regulator of transcription 1, TBP-binding (negative cofactor 2)	AB051447	1,11
KIAA1660 protein	Consensus includes gb:BF084105 /FEA=EST /DB_XREF=est:QV4-CT0491.140900-398-e06 /UG=Hs.247954 Human thrombospondin-1 gene, pa	1,11
Consensus includes gb:BF084105 /FEA=EST /DB_XREF=est:QV4-CT0491.140900-398-e06 /UG=Hs.247954 Human thrombospondin-1 gene, pa	AF135593	1,11
vacuolar protein sorting 41 (yeast)	BG289199	1,11
RNA binding motif protein 8A	BF508646	1,11
v-naf musculoaponeurotic fibrosarcoma oncogene homolog (avian)	U85943	1,11
RAE1 RNA export 1 homolog (S. pombe)	NM_006637	1,11
olfactory receptor, family 5, subfamily I, member 1	NM_004547	1,11
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15)	NM_004333	1,11
v-rif murine sarcoma viral oncogene homolog B1	AF077614	1,11
ATPase, H+ transporting lysosomal (vacuolar proton pump), member M	AF010316	1,11
prostaglandin H synthase	gb:NM_025111.1 /DEF=Homo sapiens hypothetical protein FLJ23571 (FLJ23571), mRNA, /FEA=mRNA, GEN=FLJ23571 /PROD=hypothetical protein FLJ23571 /DB_XREF=	1,11
gb:NM_025111.1 /DEF=Homo sapiens hypothetical protein FLJ23571 (FLJ23571), mRNA, /FEA=mRNA, GEN=FLJ23571 /PROD=hypothetical protein FLJ23571 /DB_XREF=	NM_014338	1,11
phosphatidylserine decarboxylase	NM_015653	1,11
DKFZP66F0548 protein	NM_014249	1,11
nuclear receptor subfamily 2, group E, member 3	NM_017609	1,11
hypothetical protein DKFZP434A1721	NM_001999	1,11
fibrillin 2 (congenital contractual arachnodactyly)	AL037534	1,11
Consensus includes gb:AL037534 /FEA=EST /DB_XREF=est:DKFZp564F2472_1 /CLONE=DKFZp564F2472 /UG=Hs.153612 ATP-binding cassette	AL050036	1,11
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	NM_014469	1,11
testes-specific heterogenous nuclear ribonucleoprotein G-T	NM_017971	1,11
mitochondrial ribosomal protein L20	NM_020241	1,11
sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B	NM_016220	1,11
zinc finger protein (ZF2D5)	AF189723	1,11
ATPase, Ca++ transporting, type 2C, member 1	NM_024664	1,11
hypothetical protein FLJ11838	D12620	1,11
cytochrome P450, subfamily IVF, polypeptide 3 (leukotriene B4 omega hydroxylase)	NM_023919	1,11
taste receptor, type 2, member 7	NM_000250	1,11
myeloperoxidase	NM_002115	1,11
hexokinase 3 (white cell)	M12959	1,11
T cell receptor alpha locus	NM_015472	1,11
transcriptional co-activator with PDZ-binding motif (TAZ)	NM_014880	1,11
KIAA0022 gene product	X74496	1,11
prolyl endopeptidase	AF263541	1,11
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 4	NM_024914	1,11
hypothetical protein FLJ13262	NM_000662	1,11
N-acetyltransferase 1 (arylamine N-acetyltransferase)	NM_002180	1,11
interleukin 17 (cytotoxic T-lymphocyte-associated serine esterase 8)	AB014576	1,11
KIAA0678 protein	NM_024712	1,11
engulfment and cell motility 3 (ced-12 homolog, C. elegans)	NM_002395	1,11
malic enzyme 1, NADP(+)-dependent, cytosolic	NM_021242	1,11
hypothetical protein STRAAT11499	U02519	1,11
collagen, type IV, alpha 3 (Goodpasture antigen)	NM_005180	1,11
murine leukemia viral (bmt-1) oncogene homolog	M13149	1,11
histidine-rich glycoprotein	NM_000238	1,11
potassium voltage-gated channel, subfamily H (eag-related), member 2	NM_014318	1,11
apoptosis related protein	NM_020125	1,11
BCL4-like membrane protein precursor	NM_024114	1,11
hypothetical protein MGC4827	AB002304	1,11
capicua homolog (Drosophila)	X07882	1,11
Consensus includes gb:X07882 /DEF=Human PRB4 gene for proline-rich protein Po, allele S /FEA=mRNA /DB_XREF=gi:35647 /UG=Hs.103972 proline-rich protein Basini ; X07882	Consensus includes gb:AK025114.1 /DEF=Homo sapiens cDNA: FLJ21481 fta, clone COL04727, /FEA=mRNA /DB_XREF=gi:10437566 /UG=Hs.306768 Homo sapiens cD AK025114	1,11
Consensus includes gb:AK025114.1 /DEF=Homo sapiens cDNA: FLJ21481 fta, clone COL04727, /FEA=mRNA /DB_XREF=gi:10437566 /UG=Hs.306768 Homo sapiens cD AK025114	NM_000286	1,11
peritonsillar biogenesis factor 12	Consensus includes gb:AK000173.1 /DEF=Homo sapiens cDNA FLJ20166 fs, clone COL09511, /FEA=mRNA /DB_XREF=gi:7020086 /UG=Hs.306386 Homo sapiens cDN AK000173	1,11
Consensus includes gb:AK000173.1 /DEF=Homo sapiens cDNA FLJ20166 fs, clone COL09511, /FEA=mRNA /DB_XREF=gi:7020086 /UG=Hs.306386 Homo sapiens cDN AK000173		

TBP-interacting protein
breast cell glutaminase
hypothetical protein FLJ13386
angiotensin vasopressin receptor 1A
FOS-like antigen-1
tubor and KH domain-containing protein
KIAA1449 protein
gbAF13007.1 /DEF=Homo sapiens clone FLB9023 PRO2425 mRNA, complete cds. /FEA=mRNA (PROD=PRO2425 /DB_XREF=gi:11493448 /UG=Hs.302148 Homo sap AF13007.1
karyopherin alpha 1 (importin alpha 5)
CD80 antigen (CD28 antigen ligand 1, B7-1 antigen)
DEADH (Asp-Glu-Ala-Asp-His) box polypeptide 28
CSJ-96 protein
dynen, axonemal, light polypeptide 4
neuron-specific protein
ADP-ribosylation factor 4-like
hypothetical protein MGC12386
hypothetical protein FLJ10330
hypothetical protein FLJ21168
cytochrome b5 outer mitochondrial membrane precursor
M27830 Human 28S ribosomal RNA gene, complete cds [5, M, 3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
truncaldehyde repeat containing 21
perlepin
putative G protein coupled receptor
Kruppel-related zinc finger protein
protease inhibitor 3, skin-derived (SKALP)
luronate 2-sulfatase (Hunter syndrome)
HRPAP20 short form
neurotensin receptor 1 (high affinity)
adaptor-related protein complex 3, beta 2 subunit
DNA-binding protein amplifying expression of surfactant protein B
estrogen receptor 2 (ER beta)
hypothetical protein MGC2748
hect domain and RLD 2
nuclear antigen Sp100
similar to rat histagral membrane glycoprotein POM121
hypothetical protein FLJ21582
Consensus includes gb:A057637 /FEA=EST /DB_XREF=gi:3331503 /DB_XREF=est:cy31h08.x1 CLONE=IMAGE:1667483 /UG=Hs.234898 ESTs, Weakly similar to 2108 A057637
mitogen-activated protein kinase 13
Consensus includes gb:AK021640.1 /DEF=Homo sapiens cDNA FLJ11578 fls, clone HEMBA1003571. /FEA=mRNA /DB_XREF=gi:10432881 /UG=Hs.287427 Homo sapie
actin, alpha 2
KIAA0725 protein
arginine-glutamic acid dipeptide (RE) repeats
NFS1 nitrogen fixation 1 (S. cerevisiae)
male-specific lethal 3-like 1 (Drosophila)
ATP/GTP-binding protein
KIAA1065 protein
small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (racalactine, neurotactin)
solute carrier family 9 (neurotransmitter transporter, taurine), member 8
KIAA0096 protein
growth arrest-specific 7
gastin
lectin, galactoside-binding, soluble, 8 (galectin 8)
target of myo1 (chicken)
phosphoribosyl pyrophosphate amidotransferase
amyloid beta precursor protein (cytoplasmic tail) binding protein 2
glutathione S-transferase A4
hypothetical protein DKFZP434I143
putative homeodomain transcription factor 1
cyclin G2
hypothetical protein FLJ10719
tyrosylprotein sulfoxidase 1
SMA4
triggering receptor expressed on myeloid cells 2

AB020836 1.11
NM_013267 1.11
NM_025180 1.11
AF030825 1.11
BG251286 1.11
NM_008862 1.11
H04342 1.11
BF575685 1.11
NM_005191 1.11
BG477502 1.11
NM_015703 1.11
AL085883 1.11
M98528 1.11
U25771 1.11
BC005311 1.11
NM_018061 1.11
BC005934 1.11
AF130113 1.11
M27830 1.11
U80756 1.11
NM_006262 1.11
NM_007223 1.11
NM_015852 1.11
L10343 1.11
A1819115 1.11
NM_020355 1.11
NM_002531 1.11
NM_004644 1.11
NM_006692 1.11
AF124790 1.11
BC001080 1.11
NM_004667 1.11
U36501 1.11
NM_014348 1.11
A128310 1.11
BC000433 1.11
AX021640 1.11
AU146889 1.11
AB018288 1.11
AF016005 1.11
NM_021100 1.11
NM_006800 1.11
NM_006831 1.11
H05666 1.11
NM_002986 1.11
U16120 1.11
A1806207 1.11
NM_005890 1.11
NM_000805 1.11
AF074000 1.11
NM_005488 1.11
U00236 1.11
NM_006380 1.11
NM_001512 1.11
AL080135 1.11
BC002447 1.11
NM_004354 1.11
W74442 1.11
NM_003596 1.11
X83300 1.11
NM_018955 1.11

hypothetical protein FLJ12820	NM_022451	1.11
Consensus includes gb:AL050074.1 IDEF+Homo sapiens mRNA: cDNA DKFZ566F1946 (from clone DKFZ566F1946). /FEA=mRNA/DB_XREF=gi:4884307 /UG=Hs.27	AK0250074	1.11
TATA box binding protein (TBP)-associated factor, RNA polymerase I, B. 63KD	AF332509	1.11
zinc finger protein 4 protein HZIC4	NM_020366	1.11
retinitis pigmentosa GTPase regulator interacting protein 1	NM_000717	1.11
carbonic anhydrase IV	AK022387	1.11
Consensus includes gb:AK022387.1 IDEF+Homo sapiens cDNA FLJ12325 fls, clone MAMMA1002125. /FEA=mRNA/DB_XREF=gi:10433772 /UG=Hs.287514 Homo sapie	NM_024944	1.11
chromosome 21 open reading frame 68	NM_004189	1.11
SRY (sex determining region Y)-box 14	D31840	1.11
dentatorubral-pallidoluysian atrophy (atrophin-1)	AI066655	1.11
SEC63 protein	T15991	1.11
cholinergic receptor, muscarinic 3	NM_025226	1.11
MSTP332 protein	AL537457	1.11
neurofilament, light polypeptide (68kD)	AL537457	1.11
Consensus includes gb:AL078636.1 IDEF+Homo sapiens mRNA (full length insert cDNA clone EUROMAGE 117929 /FEA=mRNA/DB_XREF=gi:5051866 /UG=Hs.227171	AB020690	1.11
paraneoplastic antigen MA2	AB020690	1.11
Consensus includes gb:137198.1 IDEF+Homo sapiens (clone B3B3E13) Huntingtons disease candidate region mRNA fragment. /FEA=mRNA/DB_XREF=gi:600519 /UG=	L37198	1.11
RAB17, member RAS oncogene family	NM_022449	1.11
PRO1854 protein	AF18078	1.11
8-oxoguanine DNA glycosylase	NM_016620	1.11
Consensus includes gb:BF592782 /FEA=EST/DB_XREF=gi:11685106 /DB_XREF=est:794d06.x1 /CLONE=IMAGE:3442594 /UG=Hs.289068 Homo sapiens cDNA FLJ111	BF592782	1.11
forward box G1B	NM_005249	1.11
general transcription factor IIIC, polypeptide 4 (90KD)	NM_012204	1.11
chromosome 1 open reading frame 20	T90013	1.11
hypothetical protein HT036	NM_031207	1.11
bone morphogenetic protein 1	NM_001199	1.11
spondin 1, (I-spondin) extracellular matrix protein	AB051390	1.11
G protein-coupled receptor 35	AF089087	1.11
jerky homolog (mouse)	AF072468	1.11
odorant-binding protein 2B	NM_014581	1.11
fibroblast growth factor 3 (murine mammary tumor virus integration site (v-int-2) oncogene homolog)	NM_005247	1.11
six transmembrane epithelial antigen of the prostate	NM_012449	1.11
interleukin 1 receptor-like 1	NM_003856	1.11
chemokine (C-C motif) receptor 4	NM_005508	1.11
cathepsin Z	AF073890	1.11
maternally expressed 3	AF090934	1.11
Thy-1 cell surface antigen	AL161958	1.11
putative DNA binding protein	AL117354	1.11
Consensus includes gb:BC389837 /FEA=EST/DB_XREF=gi:13292385 /DB_XREF=est:802440589F1 /CLONE=IMAGE:4551478 /UG=Hs.289056 ESTs, Highly similar to 1: BG398837	BC389837	1.10
Consensus includes gb:AW814026 /FEA=EST/DB_XREF=gi:7907020 /DB_XREF=est:RC6-ST0198-071199-021-G02 /UG=Hs.273621 Homo sapiens cDNA: FLJ21350 fls, AW814026	NM_005123	1.10
nuclear receptor subfamily 1, group H, member 4	M22921	1.10
UDP-Gal:beta-GlcNAc beta 1,4- galactosyltransferase, polypeptide 1	NM_007252	1.10
Retina-derived POU-domain factor-1	AF112345	1.10
Integrin, alpha 10	NM_022720	1.10
DiGeorge syndrome critical region gene 8	NM_005885	1.10
golgi autoantigen, golgin subfamily a, 3	NM_002138	1.10
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	NM_007098	1.10
clathrin, heavy polypeptide-like 1	AW511135	1.10
nudix (nucleoside diphosphate linked moiety X)-type motif 4	NM_005160	1.10
adennergic, beta, receptor kinase 2	AW134976	1.10
KIAA0984 protein	BC005111	1.10
glutamate dehydrogenase 1	NM_002781	1.10
pregnancy specific beta-1-glycoprotein 5	NM_016355	1.10
hsp256 protein	AL041124	1.10
hypothetical protein PP1665	NM_025040	1.10
hypothetical protein FLJ21841	NM_183533	1.10
Consensus includes gb:AL163533 /DEF=Human DNA sequence from clone RP11-114G1 on chromosome 13q31.3-33.1 Contains a pseudogene similar to transcription fac	AL163533	1.10
acyl-Coenzyme A oxidase 1, palmitoyl	S68189	1.10
interferon, alpha 14	NM_002172	1.10
protein inhibitor of activated STAT protein PIASy	AK022461	1.10
apoptosis regulator BCL-G	NM_030766	1.10
mitogen-activated protein kinase-activated protein kinase 2	BG325734	1.10
protein tyrosine phosphatase, receptor type, I polypeptide (PTPRF), interacting protein (Iiprin), alpha 3	BE501428	1.10

hypothetical protein FLJ12476
 sulfolipase, estrogen-prefering
 Consensus includes gb:AK024158.1 /DEF=Homo sapiens cDNA FLJ14098 fta, clone MAMMA1000752. /FEA=mRNA /DB_XREF=gi:10436469 /UG=Hs.303289 Homo sapie
 frizzled homolog 4 (Drosophila)
 GN3 general control of amino-acid synthesis 5-like 1 (yeast)
 synaptob, gamma 1
 Ac-like transposable element
 endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2
 fasciculation and elongation protein zeta 2 (zyglin II)
 amyloid P component, serum
 angiotensin receptor 2
 PRO0481 protein
 hypothetical protein FLJ10520
 Cdc42 effector protein 4; binder of Rho GTPases 4
 KIAA0263 protein
 Consensus includes gb:U92706 /DEF=Human rearranged immunoglobulin heavy chain (A1VH3) gene, partial cds /FEA=CDS /DB_XREF=gi:2155271 /UG=Hs.249245 Hom
 MORF-related gene X
 hypothetical protein FLJ13105
 phosphatidylinositol glycan, class B
 Consensus includes gb:N32526 /FEA=EST /DB_XREF=gi:1152925 /DB_XREF=est:Y1104.s1 /CLONE=IMAGE:270943 /UG=Hs.300642 serologically defined colon cancer
 tumor endothelial marker 5 precursor
 endostatin alpha
 HIRA interacting protein 5
 zinc finger protein 181 homolog (mouse)
 spectrin, beta, non-erythrocytic 2
 ephrin-A5
 hypothetical protein FLJ20085
 Consensus includes gb:AF090887.1 /DEF=Homo sapiens clone HQ0085. /FEA=mRNA /DB_XREF=gi:6690154 /UG=Hs.306562 Homo sapiens clone HQ0085
 transient receptor potential channel 4
 1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)
 COP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2
 hypothetical protein FLJ20113
 Consensus includes gb:AK024173.1 /DEF=Homo sapiens cDNA FLJ14111 fta, clone MAMMA1001630. /FEA=mRNA /DB_XREF=gi:10436489 /UG=Hs.288567 Homo sapie
 Consensus includes gb:AK024123.1 /DEF=Homo sapiens cDNA FLJ14061 fta, clone HEMBB1000748. /FEA=mRNA /DB_XREF=gi:10436427 /UG=Hs.306664 Homo sapie
 protease, serine, 4 (trypsin 4, brain)
 chromosome 10 open reading frame 3
 C-terminal binding protein 2
 hairless homolog (mouse)
 hypothetical protein FLJ20245
 endonuclease G-like 2
 FYN oncogene related to SRC, FGR, YES
 endothelial PAS domain protein 1
 PRO1578 protein
 TAP1 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 250 kD
 gonadotropin-releasing hormone 1 (luteinizing-releasing hormone)
 cathepsin F
 DKFZP586K0524 protein
 Opa-interacting protein 5
 solute carrier family 19, member 3
 thyroid peroxidase
 carboxypeptidase E
 sodium channel, voltage-gated, type I, beta polypeptide
 origin recognition complex, subunit 6 (yeast homolog)-like
 hypothetical protein FLJ10921
 myosin IF
 T-box 2
 kinesin 1 (kinesin receptor)
 DKFZP584F1422 protein
 hypothetical protein FLJ23519
 ribosomal protein S6 kinase, 90kD, polypeptide 4
 chromosome 11 open reading frame 8
 K01391 B subtilis TrpE protein, TrpC protein corresponding to nucleotides 1883-4400 of K01391 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 K01391)

1.10 NM_022784
 1.10 NM_005420
 1.10 AK024158
 1.10 NM_012193
 1.10 NM_004492
 1.10 NM_001487
 1.10 NM_018987
 1.10 NM_004729
 1.10 NM_001401
 1.10 AL117593
 1.10 NM_001639
 1.10 U16937
 1.10 NM_031288
 1.10 BC002574
 1.10 AL162074
 1.10 AB008625
 1.10 U92706
 1.10 NM_012286
 1.10 NM_025001
 1.10 AU144243
 1.10 N32526
 1.10 BF511315
 1.10 AF157510
 1.10 NM_015700
 1.10 AL534416
 1.10 NM_006946
 1.10 NM_001962
 1.10 NM_017660
 1.10 AF090887
 1.10 NM_016179
 1.10 U56418
 1.10 Y16521
 1.10 NM_017870
 1.10 AK024173
 1.10 AK024123
 1.10 AW007273
 1.10 NM_001214
 1.10 N23018
 1.10 NM_018411
 1.10 NM_017723
 1.10 NM_005728
 1.10 N20923
 1.10 AF052094
 1.10 AF118067
 1.10 X07024
 1.10 NM_000825
 1.10 NM_003793
 1.10 AL050040
 1.10 BE045993
 1.10 NM_025243
 1.10 M17755
 1.10 AB22855
 1.10 NM_001037
 1.10 NM_014321
 1.10 NM_018272
 1.10 BF740152
 1.10 NM_005994
 1.10 NM_004986
 1.10 AL049942
 1.10 AK027172
 1.10 NM_003942
 1.10 BE645386
 1.10 K01391

Consensus includes gb:AF065241.1 /DEF=Homo sapiens thioredoxin delta 3 (TXN delta 3) mRNA, partial cds. /FEA=mRNA /GEN=TXN delta 3 /PROD=thioredoxin delta 3 AF065241 1.10

Dicoforge syndrome gene H L77565 1.10

fragile X mental retardation 2 L75569 1.10

lymphoid blast crisis oncogene BF222823 1.10

ATPase, H⁺/K⁺ transporting, nongastric, alpha polypeptide NM_001676 1.10

gb:BC005914.1 /DEF=Homo sapiens, clon MGC:14510, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:14510) /DB_XREF=gi:13543511 /FL=gb:BC005914 1.10

CDC2-related protein kinase 7 NM_016507 1.10

Fanconi anemia, complementation group A A983201 1.10

harakuri, BCL2 interacting protein (contains only BH3 domain) U76376 1.10

insulin-like growth factor 2 receptor NM_008876 1.10

HZF-w1 protein NM_025236 1.10

tuberous sclerosis 1 AF013168 1.10

serum response factor (c-fos serum response element-binding transcription factor) A1188786 1.10

hypothetical protein FLJ20265 NM_017733 1.10

zinc finger protein 84 (HPF2) NM_003428 1.10

TNF receptor-associated factor 6 NM_004620 1.10

pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1) M57389 1.10

Consensus includes gb:A1346181 /FEA=EST /DB_XREF=gi:4083387 /DB_XREF=est:qp49603.x1 /CLONE=IMAGE:1928317 /UG=Hs.42712 MAX protein A1346181 1.10

DIPB protein AA176780 1.10

Epha2 NM_004431 1.10

hypothetical protein FLJ13195 similar to stromal antigen 3 NM_022906 1.10

corbotropin releasing hormone receptor 1 NM_004382 1.10

hypothetical protein FLJ13116 NM_016388 1.10

Kruppel-like factor 13 NM_015995 1.10

protein kinase, CAMP-dependent, regulatory, type I, beta A014660 1.10

hypothetical protein FLJ14251 AA504646 1.10

meningioma (disrupted in balanced translocation) 1 NM_002430 1.10

likely ortholog of mouse actin-related protein 8 homolog (S. cerevisiae) NM_022899 1.10

retinoblastoma binding protein 2 NM_005056 1.10

choroideremia (Rab escort protein 1) NM_000350 1.10

TAR DNA binding protein NG111668 1.10

consensus includes gb:AU146983 /FEA=EST /DB_XREF=gi:11008504 /DB_XREF=est:AU146983 /CLONE=HEMBB1002049 /UG=Hs.262212 Homo sapiens cDNA FLJ121 AU146983 1.10

consensus includes gb:AU146983 /FEA=EST /DB_XREF=gi:11008504 /DB_XREF=est:AU146983 /CLONE=HEMBB1002049 /UG=Hs.262212 Homo sapiens cDNA FLJ121 AU146983 1.10

Homer, neuronal immediate early gene, 2 Y19026 1.10

diacylglycerol kinase, epsilon (6A0D) NM_003647 1.10

Bacillus subtilis /REF=M24537 /DEF=B subtilis pheA, pheA genes corresponding to nucleotides 2484-2875 of M24537 /LEN=1409 (-5, -M, -3 represent transcript regions 5 M24537 1.10

Consensus includes gb:AL021327 /DEF=Human DNA sequence from PAC 12409 on chromosome 8q21. Contains DNAJ2 (HduJ) like pseudogene, ESTs, STSs and GSSs AL021327 1.10

ADP-ribosyltransferase (NAD⁺: poly (ADP-ribose) polymerase)-like 3 AF083068 1.10

Consensus includes gb:AV727101 /FEA=EST /DB_XREF=gi:10836522 /DB_XREF=est:AV727101 /CLONE=HTC8GF05 /UG=Hs.25272 E1A binding protein p300 /FL=gb:AV727101 1.10

HSPC059 protein NM_016536 1.10

growth factor, augmenter of liver regeneration (ERV1 homolog, S. cerevisiae) NM_005262 1.10

carboxypeptidase A4 NM_016352 1.10

sorting nexin 4 NM_003794 1.10

retinoid acid receptor responder (tazarotene induced) 2 BC000069 1.10

Consensus includes gb:AF164963.1 /DEF=Homo sapiens tumor antigen NA88-A pseudogene, complete sequence. /FEA=mRNA /DB_XREF=gi:5901726 /UG=Hs.306576 1.10

Rh type C glycoprotein NM_016321 1.10

Consensus includes gb:U86813.1 /DEF=Homo sapiens serotonin-7 receptor pseudogene, complete sequence. /FEA=mRNA /PROD=serotonin-7 receptor /DB_XREF=gi:31 U86813 1.10

insulin receptor substrate 2 BF700086 1.10

Sec23 homolog B (S. cerevisiae) BC005032 1.10

a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 2 NM_014244 1.10

putative secreted protein NM_015926 1.10

T-cell lymphoma invasion and metastasis 1 NM_003253 1.10

Consensus includes gb:AF195953 /DEF=Homo sapiens membrane-bound aminopeptidase P (XNPEP2) gene, complete cds /FEA=mRNA /DB_XREF=gi:11066156 /UG=Hs. AF195953 1.10

visinin-like 1 AF039555 1.10

hypoxia-inducible protein 2 NM_013332 1.10

hypothetical protein FLJ10103 NM_017996 1.10

chorionic somatomammotropin hormone 2 NM_022844 1.10

HYA22 protein AA129726 1.10

nuclear factor I/X (CCAAT-binding transcription factor) U18759 1.10

HBV pX associated protein-8 NM_016578 1.10

KUAI1069 protein AW665865 1.10

Consensus includes gb:AW971983 /FEA=EST /DB_XREF=gi:8161949 /DB_XREF=est:EST384192 /UG=Hs.293003 ESTs, Moderately similar to ALU4_HUMAN ALU SUBF AW971983 1.10

NGF1A binding protein 2 (EGR1 binding protein 2)
 gb:NM_024907.1 /DEF=Homo sapiens hypothetical protein FLJ11798 (FLJ11798), mRNA, /FEA=mRNA /GEN=FLJ11798 /PROD=hypothetical protein FLJ11798 /DB_XREF=NM_024907
 hypothetical protein FLJ22529
 hepcidin-associated membrane protein 1
 neuropilin 1
 immunoglobulin kappa variable 1D-8
 zinc finger protein 278
 N-myristoyltransferase 1
 ubiquitin specific protease 22
 DESC1 protein
 phosphatase, glycogen; liver (Hers disease, glycogen storage disease type VI)
 chromosome 22 open reading frame 4
 human T-cell leukemia virus enhancer factor
 nucleoside diphosphate kinase type 6 (inhibitor of p53-induced apoptosis-alpha)
 chloride channel 5 (nephrolithiasis 2, X-linked; Dent disease)
 KIAA0871 protein
 Chediak-Higashi syndrome 1
 transcription factor CP2
 progastrin (pepsinogen C)
 neurofibromin 2 (bilateral acoustic neuroma)
 methyl CpG binding protein 2 (Rett syndrome)
 cytochrome P450, 51 (lanosterol 14-alpha-demethylase)
 hypothetical protein FLJ10420
 septaplerin reductase (7,8-dihydrobiopterin:NADP+ oxidoreductase)
 deiodinase, iodothyronine, type II
 splicing factor (45KD)
 sec13-like protein
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9
 general transcription factor IIE, polypeptide 1 (alpha subunit, 56kD)
 hypothetical protein MGC11061
 forhead box D1
 KIAA0869 protein
 RNA binding motif protein 7
 slug homolog, zinc finger protein (chicken)
 mitochondrial ribosomal protein 63
 cyclin-dependent kinase 5, regulatory subunit 2 (p39)
 KIAA0749 protein
 hypothetical protein FLJ20828
 translin-associated factor X
 membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125)
 prostaglandin I2 (prostaglandin) receptor (IP)
 hypothetical protein FLJ23306
 Consensus includes gb:A093572 /FEA=EST /DB_XREF=gi:3432548 /DB_XREF=est:gb1511.x1 /CLONE=IMAGE:1686365 /UG=Hs.18081 Homo sapiens clone DT1P186
 mannosyl (beta-1,4)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase
 hypothetical protein FLJ12875
 stratifin
 ubiquitin-conjugating enzyme E2L 3
 hypothetical protein FLJ13659
 Consensus includes gb:AL109709.2 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 43432, /FEA=mRNA /DB_XREF=gi:9187596 /UG=Hs.167456
 diacylglycerol kinase, gamma (90kD)
 synovial sarcoma, X breakpoint 5
 DKFZP564D116 protein
 Sp2 transcription factor
 cell division cycle 42 (GTP binding protein, 25kD)
 protein phosphatase, EF hand calcium-binding domain 1
 hypothetical protein FLJ21069
 growth arrest-specific 11
 adaptor-related protein complex 4, epsilon 1 subunit
 ribosomal protein L28
 DAZ associated protein 2
 YY1-associated factor 2
 chloride channel 4
 mitogen-activated protein kinase kinase 4

capping protein (actin filament) muscle Z-line, alpha 1
 pregnancy-associated plasma protein-E
 protease, serine, 11 (IGF binding)
 Cdc42 effector protein 3
 Fanconi anemia, complementation group A
 hypothetical protein FLJ10850
 PAP-1 binding protein
 synaptophysin 4
 KIAA0972 protein
 arginine vasopressin receptor 1A
 DKFZP564O0423 protein
 REV3-like, catalytic subunit of DNA polymerase zeta (yeast)
 tricholein repeat containing 1
 annexin A11
 signal transducer and activator of transcription 1, 91 kD
 zinc finger protein 278
 CDC14 cell division cycle 14 homolog A (S. cerevisiae)
 neutralized-like (Drosophila)
 Consensus includes gb:BE794663 /FEA=EST /DB_XREF=gi:10215861 /DB_XREF=est:601590459F1 /CLONE=IMAGE:3944463 /UG=Hs.306688 Homo sapiens cDNA FLJ BE794663
 GLI-Kruppel family member GLI2
 gamma-glutamyl hydrolase (conjugase, poly(glycylglycyl) hydrolase)
 interleukin 6 signal transducer (gp130, oncostatin M receptor)
 Consensus includes gb:AK022254.1 /DEF=Homo sapiens cDNA FLJ12192 lis, clone MAMMA1000851, /FEA=mRNA /DB_XREF=gi:10433612 /UG=Hs.288451 Homo sapiens cDNA FLJ BE794663
 complement component 1, q subcomponent, receptor 1
 paired mesoderm homeobox 2b
 Cluster Ind. A472320:jb7c02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:2148482 /clone_end=3 /gb=A1472320 /gi=4334410 /ug=Hs.48504 /len=548
 smoothelin
 C-terminal binding protein 2
 PRO0297 protein
 centaurin, delta 2
 hypothetical protein DKFZp586E1923
 myocilin, trabecular meshwork inducible glucocorticoid response
 hypothetical protein FLJ14220
 Consensus includes gb:AL035308.1 /DEF=H. sapiens gene from PAC 42816, /FEA=CDS /PROD=hypothetical protein /DB_XREF=gi:4200237 /UG=Hs.105911 H. sapiens ge
 homolog of mouse quaking QKI (KH domain RNA binding protein)
 ankrym 3, node of Ranvier (ankyrin G)
 hypothetical protein FLJ10210
 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2
 natriuretic peptide receptor Aguanyle cyclase A (atrionatriuretic peptide receptor A)
 aminolevulinatase, delta-, dehydratase
 core-binding factor, runt domain, alpha subunit 2; translocated to, 2
 FYN oncogene related to SRC, FGR, YES
 hypothetical protein FLJ2242
 neuronal protein
 hypothetical protein FLJ12455
 Consensus includes gb:AW469184 /FEA=EST /DB_XREF=gi:7039290 /DB_XREF=est:hc78gp4.x1 /CLONE=IMAGE:2898870 /UG=Hs.65406 ESTs
 Consensus includes gb:AL137325.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M0835 (from clone DKFZp434M0835), /FEA=mRNA /DB_XREF=gi:6807816 /UG=Hs.27
 small inducible cytokine subfamily A (Cys-Cys), member 15
 crystallin, beta B3
 galanin-related peptide
 aldehyde dehydrogenase 7 family, member A1
 chromodomain helicase DNA binding protein 2
 Consensus includes gb:U05589.1 /DEF=Human ribosomal protein S1 homolog mRNA, partial cds, /FEA=mRNA /DB_XREF=gi:497001 /UG=Hs.371 ribosomal protein S1-III
 gb:U72513.1 /DEF=Human RPL13-2 pseudogene mRNA, complete cds, /FEA=mRNA /PROD=RPL13-2 /DB_XREF=gi:1673516 /UG=Hs.159486 Human RPL13-2 pseudog
 Kruppel-like factor 5 (intestinal)
 nuclear protein, axilla-telangiectasia locus
 schaefer-scute complex-like 1 (Drosophila)
 POU domain, class 5, transcription factor 1
 CD1E antigen, e polypeptide
 zinc finger protein 192
 collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase
 Consensus includes gb:A1279819 /FEA=EST /DB_XREF=gi:3918053 /DB_XREF=est:qm26h04.x1 /CLONE=IMAGE:1882999 /UG=Hs.27373 Homo sapiens mRNA; cDNA [A1279819

277

corticotropin releasing hormone binding protein
 Consensus includes gb:AL122093.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B2115 (from clone DKFZp434B2115), /FEA=mRNA /DB_XREF=gi:6102904 /UG=Hs.13 /AL122093
 guanylate cyclase 2D, membrane (retina-specific)
 eukaryotic translation initiation factor 4E binding protein 2
 small inducible cytokine subfamily A (Cys-Cys), member 8 (monocyte chemotactic protein 2)
 corticotropin releasing hormone receptor 2
 contactin associated protein 1
 pili-like transcription factor
 chordin
 prothymosin-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II
 Consensus includes gb:T15769 /FEA=EST /DB_XREF=gi:517828 /DB_XREF=est:181859 /UG=Hs.321572 Homo sapiens clone 25048 mRNA sequence
 glioblastoma amplified sequence
 adenylate cyclase activating polypeptide 1 (pituitary) receptor type I
 monoglyceride lipase
 ephrin-A1
 deiodinase, liothyronine, type II
 translation initiation factor IF2
 C3HC4-like zinc finger protein
 KIAA1067 protein
 PRKc, apoptosis, WT1, regulator
 5-hydroxytryptamine (serotonin) receptor 2B
 peroxisomal biogenesis factor 11A
 zinc finger protein 272
 gb:AF049656.1 /DEF=Homo sapiens Inducible nitric oxide synthase (NOS) mRNA, complete cds. /FEA=CDS /GEN=INOS /PROD=inducible nitric oxide synthase /DB_XREF=AF049656
 Cluster incl. A069655, wc1205, x1 Homo sapiens cDNA, 3' end /done=IMAGE:2314928 /done_end=3 /gb=A069655 /gi=4834429 /ug=Hs.30251 /len=649
 Consensus includes gb:AL121948 /FEA=Human DNA sequence from clone RP3-398E4 on chromosome 6. Contains the 3' end of the SIM1 gene for single-minded (Drosop
 Consensus includes gb:BC536224 /FEA=EST /DB_XREF=gi:13527769 /DB_XREF=est:60256445F1 /CLONE=IMAGE:4690258 /UG=Hs.123030 Human kappa-immunoglobulin BG536224
 gb:AF090900.1 /DEF=Homo sapiens clone HQ0189 PRO0189 mRNA, complete cds. /FEA=mRNA /PROD=PRO0189 /DB_XREF=gi:6690176 /UG=Hs.91393 Homo sapien: AF090900
 W22117
 HT017 protein
 hypothetical protein FLJ20699
 endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 4
 p53 target zinc finger protein
 amiloride binding protein 1 (amine oxidase (copper-containing))
 kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
 TNF receptor-associated factor 5
 gb:NM_019398.1 /DEF=Homo sapiens kallikrein 12 (KLK12), mRNA. /FEA=mRNA /GEN=KLK12 /PROD=kallikrein 12 /DB_XREF=gi:9665235 /UG=Hs.159679 kallikrein 12 /NM_019398
 nuclear receptor subfamily 2, group F, member 2
 hypothetical protein PRO1331
 hypothetical protein FLJ20396
 putative Rab5 GDP/GTP exchange factor homologue
 KIAA0995 protein
 carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5
 carbodiimide 12
 peptidylprolyl isomerase E (cyclophilin E)
 lymphoid blast crisis oncogene
 claudin 9
 hypothetical protein FLJ22059
 DKFZP386N0721 protein
 myeloid/lymphoid or mixed-lineage leukemia (trithorax homologue, Drosophila), translocated to, 2
 dynamin 2
 hypothetical protein MGC4771
 hypothetical protein FLJ10647
 mannosidase, alpha, class 1A, member 1
 ras responsive element binding protein 1
 phosphodiesterase 6B, cGMP-specific, rod, beta (congenital stationary night blindness 3, autosomal dominant)
 ciliary neurotrophic factor
 elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2
 erythrocyte membrane protein band 4.1-like 3
 histone acetyltransferase
 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7)
 caveolin 3
 thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)
 chondroitin 4-sulfotransferase

T cell receptor alpha locus	AE000659	1.09
phosphatidylinositol glycan, class L	AI742901	1.09
slit homolog 2 (Drosophila)	AF055585	1.09
hypothetical protein FLJ2393	NM_025106	1.09
heat shock 90kD protein 1, alpha	BG420237	1.09
hypothetical protein FKS028	W28828	1.09
deoxyribonuclease I	M55983	1.09
glycerol kinase	X88285	1.09
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 6	NM_004775	1.09
mutL homolog 3 (E. coli)	NM_014381	1.09
sterile alpha and HEAT/Armadillo motif protein, ortholog of Drosophila	AJ280445	1.09
Consensus includes gb:AW593996 /FEA=EST /DB_XREF=gi:7281254 /DB_XREF=est:hg41g06.x1 /CLONE=IMAGE:2948218 /UG=Hs:124969 Homo sapiens clone 24707	AW593996	1.09
KIAA0687 protein	AI978623	1.09
KIAA0435 gene product	AB007895	1.09
KIAA0674 protein	AB014574	1.09
Consensus includes gb:AF090883.1 /DEF=Homo sapiens clone HQ0021. /FEA=mRNA /DB_XREF=gi:6690149 /UG=Hs:306560 Homo sapiens clone HQ0021	AF090883	1.09
nasopharyngeal carcinoma susceptibility protein	NM_013275	1.09
DKFZP564B167 protein	NM_015415	1.09
cornedoesmosin	NM_001264	1.09
MORF-related gene 15	NM_006791	1.09
Consensus includes gb:AL031600 /DEF=Human DNA sequence from clone 390E6 on chromosome 18. Contains the 3 part of the CLCN7 gene for chloride channel 7. EST: LQ3380	NM_003160	1.09
gonadotropin-releasing hormone receptor	AL080137	1.09
DKFZP434J183 protein	NM_012101	1.09
tripartite motif-containing 28	AI984421	1.09
GpE-like protein co-chaperone	AF229082	1.09
death effector filament-forming Ced-4-like apoptosis protein	NM_017612	1.09
hypothetical protein DKFZP434E2220	AL049449	1.09
Consensus includes gb:AL049449.1 /DEF=Homo sapiens mRNA: cDNA DKFZP586B1722 (from clone DKFZP586B1722) /FEA=mRNA /DB_XREF=gi:4500234 /UG=Hs:80	AL049449	1.09
KIAA0805 protein	AI815033	1.09
centromere protein F (350/400kD, mitotin)	U30872	1.09
nucleolar protein 1 (120kD)	NM_006170	1.09
chromatin assembly factor 1, subunit A (p150)	BF000239	1.09
ephA-B3	NM_001406	1.09
brain-specific angiogenesis inhibitor 2	NM_001703	1.09
inhibitor of Brulon's tyrosine kinase	AF235049	1.09
ectodermal-neural crest (with BTB-like domain)	NM_003633	1.09
PRO0811 protein	NM_014076	1.09
insulin-like growth factor 2 (somatomedin A)	M17663	1.09
oculocutaneous albinism II (pink-eye dilution (murine) homolog)	NM_000275	1.09
core promoter element binding protein	U51869	1.09
perflavin	NM_002666	1.09
alpha-methylacyl-CoA racemase	AA888589	1.09
keratin, hair, basic, 4	Y19209	1.09
Consensus includes gb:AL137715.1 /DEF=Homo sapiens mRNA: cDNA DKFZP434K0423 (from clone DKFZP434K0423); partial cds. /FEA=mRNA /GEN=DKFZP434K0423	AL137715	1.09
ribosomal protein L21	AL356414	1.09
FtsJ homolog 1 (E. coli)	NM_012280	1.09
extracellular matrix protein 1	U85932	1.09
microtubule-associated protein like echinoderm EMAP	NM_012155	1.09
cell division cycle 25A	AI343459	1.09
hypothetical protein FLJ20298	NM_017752	1.09
amyloid beta (A4) precursor protein-binding, family A, member 2 (X11-like)	AB014719	1.09
proline-serine-threonine phosphatase interacting protein 2	NM_024430	1.09
alpha-fetoprotein	NM_001134	1.09
transcription factor AP-2 alpha (activating enhancer binding protein 2 alpha)	NM_003220	1.09
RAD52 homolog (S. cerevisiae)	AF125949	1.09
proline-rich protein BstNI subfamily 1	NM_005039	1.09
casepase 4, apoptosis-related cysteine protease	U25804	1.09
calcitonin, D1 dopamine receptor-interacting protein	NM_015722	1.09
folistatin-like 3 (secreted glycoprotein)	NM_005680	1.09
kinesin-like 1	NM_004523	1.09
CGI-111 protein	NM_016048	1.09
Consensus includes gb:AC004490 /DEF=Homo sapiens chromosome 19, cosmid R29381 /FEA=CDS /DB_XREF=gi:2995576 /UG=Hs:156015 Homo sapiens chromosome AC004490	AC004490	1.09
chromosome 1 open reading frame 22	NM_025191	1.09

DKF2P564M082 protein	NM_014042	1.09
intrastemal A particle-promoted polypeptide	NM_005897	1.09
ribosomal protein L17	AL049597	1.09
DNA-crosslink repair gene SNM1	D42045	1.09
ubiquitin-conjugating enzyme E2B (RAD6 homolog)	R99037	1.09
sedmin-like 2	A4877765	1.09
regulatory factor X-associated protein	NM_007029	1.09
Consensus includes gb:U79248.1 /DEF=Human done 23826 mRNA sequence, /FEA=mRNA /DB_XREF=gi:1710195 /UG=Hs.12484 Human done 23826 mRNA sequence	NM_000538	1.09
deltatubulin	U78248	1.09
delta-tubulin	NM_030932	1.09
HCGIV-6 protein	BC000258	1.09
Consensus includes gb:AJ307760 /FEA=EST /DB_XREF=gi:4002364 /DB_XREF=eat:ib24p09.x1 /CLONE=IMAGE:2055328 /UG=Hs.14074 Homo sapiens clone 24627 mR	AF036973	1.09
S100 calcium binding protein A8 (calgranulin A)	NM_002964	1.09
glycophorin A (includes MN blood group)	A1240545	1.09
TAF-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD	BF058726	1.09
hypothetical protein FLJ20281	NM_017748	1.09
Consensus includes gb:AW341200 /FEA=EST /DB_XREF=gi:6837826 /DB_XREF=est:xx295g12.x1 /CLONE=IMAGE:2872006 /UG=Hs.186882 Homo sapiens cDNA: FLJ21	AW341200	1.09
adaptor-related protein complex 3, delta 1 subunit	A424923	1.09
UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter	N80922	1.09
serine palmitoyltransferase, long chain base subunit 2	U15555	1.09
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1	AF119873	1.09
claudin 15	NM_014343	1.09
phospholipase C, beta 2	BC000939	1.09
cholinergic receptor, nicotinic, alpha polypeptide 6	NM_004198	1.09
myeloid leukemia factor 1	NM_022443	1.09
plasmalemma vesicle associated protein	AF326591	1.09
H328	NM_015726	1.09
isovaleryl Coenzyme A dehydrogenase	M34192	1.09
regulatory factor X, 3 (influences HLA class II expression)	NM_002919	1.09
homolog of yeast MOG1	NM_014185	1.09
Consensus includes gb:AK026238.1 /DEF=Homo sapiens cDNA: FLJ22585 fta, clone HSI02678, /FEA=mRNA /DB_XREF=gi:10439027 /UG=Hs.287701 Homo sapiens cD	AK026238	1.09
Consensus includes gb:AL136318 /DEF=Human DNA sequence from clone RP1-254P11 on chromosome 10 Contains ESTs, STSs and GSSs. Contains a gene for a novel	AL136318	1.09
adducin 1 (alpha)	AL556041	1.09
protocadherin 11	NM_014522	1.09
nucleosome assembly protein 1-like 2	NM_021863	1.09
T cell receptor alpha locus	AW550865	1.09
RCD1 required for cell differentiation1 homolog (S. pombe)	A1567462	1.09
protein x 0004	NM_016301	1.09
prostate epithelium-specific Ets transcription factor	A1745526	1.09
chondroitin sulfate proteoglycan 2 (versican)	BF590263	1.09
inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	AL022726	1.09
acrosomal vesicle protein 1	NM_020114	1.09
hypothetical protein FLJ22686	NM_024957	1.09
low density lipoprotein receptor (familial hypercholesterolemia)	A861942	1.09
selectin P (granule membrane protein 140kD, antigen CD62)	NM_003005	1.09
hypothetical protein FLJ20280	NM_017741	1.09
ficollin (collagen/vitronectin domain-containing) 3 (Hakata antigen)	NM_003665	1.09
interferon regulatory factor 5	AI028035	1.09
death effector filament-forming Ced-4-like apoptosis protein	AF229061	1.09
U14573 Human Alu-Sq subfamily consensus sequence,	U14573	1.09
related RAS viral (v-ras) oncogene homolog	NM_006270	1.09
teratin	AF116810	1.09
Rho guanine nucleotide exchange factor (GEF) 3	NM_019535	1.09
ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)	BC004247	1.09
microtubule-associated protein 7	AJ242502	1.09
DKF2P34O047 protein	NM_015594	1.09
decorin	AF138302	1.09
chromosome 16 open reading frame 7	NM_004913	1.09
KUAA0825 protein	NM_015046	1.09
neuregulin 1	NM_013957	1.09
hypothetical protein PRO0388	NM_018562	1.09
Consensus includes gb:AU147598 /FEA=EST /DB_XREF=gi:11009119 /DB_XREF=est:AU147598 /CLONE=MAMMA1001143 /UG=Hs.296707 Homo sapiens cDNA FLJ12	AU147598	1.09

microtubule-associated protein, RP/EB family, member 2	AB016823	1.09
goosecoid-like	NM_005315	1.09
hypothetical protein FLJ23468	NM_024629	1.09
KIAA1026 protein	AB015329	1.09
KIAA1088 protein	AB029012	1.09
LentC (bacterial lantibiotic synthetase component C)-like 1	AB352555	1.09
hypothetical protein FLJ21551	NM_024801	1.09
hypothetical protein FLJ10110	NM_017998	1.09
Machado-Joseph disease (spinocerebellar ataxia 3, olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3)	AB68099	1.09
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 4 (GalNAc-T4)	NM_003774	1.09
Siegrin syndrome antigen A2 (60kD, ribonucleoprotein autoantigen SS-A/Ro)	AL538601	1.09
hypothetical protein FLJ21140	NM_024778	1.09
proteasome (prosome, macropain) 26S subunit, non-ATPase, 9	BC002383	1.09
zinc finger protein 274	NM_016325	1.09
hypothetical protein FLJ20241	AW163030	1.09
carboxypeptidase N, polypeptide 1, 50kD	NM_001308	1.09
desmocollin 3	NM_001941	1.09
prosapin (variant Gaucher disease and variant metachromatic leukodystrophy)	M32221	1.09
rRNA isopentenylpyrophosphate transferase	NM_017846	1.09
ras inhibitor	NM_004292	1.09
olfactory receptor, family 2, subfamily F, member 1	NM_012369	1.09
Consensus includes gb:L23852.1 /DEF=Homo sapiens (clone Z146) retinal mRNA, 3 end and repeat region /FEA=mRNA /DB_XREF=gi:393126 /UG=Hs.73838 Homo sapi	L23852	1.09
stress 70 protein chaperone, microsome-associated, 60kD	NM_006948	1.09
glutamyl aminopeptidase (aminopeptidase A)	L12468	1.09
death effector domain-containing	NM_004216	1.09
solute carrier family 1 (glial high affinity glutamate transporter), member 3	NM_004172	1.09
complement component 1, q subcomponent, receptor 1	NM_012072	1.09
B-cell CLL/lymphoma 11A (zinc finger protein)	NM_018014	1.09
hypothetical protein MGC1136	NM_024025	1.09
eukaryotic translation initiation factor 1A	BE542684	1.09
FXYD domain-containing ion transport regulator 3	NM_005971	1.09
nuclear ribotic apparatus protein 1	NM_006185	1.09
dynactin 1 (p150, glued homolog, Drosophila)	BC006163	1.09
liver-specific bHLH-Zip transcription factor	NM_015925	1.09
OC6 protein	NM_020189	1.09
KIAA1639 protein	AB051446	1.09
hypothetical protein FLJ23438	NM_024671	1.09
collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive)	L23982	1.09
Consensus includes gb:AI432713 /FEA=EST /DB_XREF=gi:4283899 /DB_XREF=est:th3e02.x1 /CLONE=IMAGE:2121050 /UG=Hs.241053 ESTs	AI432713	1.09
DNA segment on chromosome X and Y (unique) 155 expressed sequence	M99578	1.09
cannabinoid receptor 1 (brain)	U73304	1.09
alpha thalassemia/mental retardation syndrome X-linked (RAO54 homolog, S. cerevisiae)	AB50257	1.09
lipoprotein lipase	AB479175	1.09
H2.O-like homeo box 1 (Drosophila)	BF872975	1.09
mannosidase, alpha, class 1A, member 2	M60721	1.09
likely ortholog of mouse zinc finger homeodomain 4	H97840	1.09
regulated in glioma	NM_024721	1.09
GDP-mannose 4,6-dehydratase	NM_006394	1.09
hypothetical protein FLJ13322	A1762113	1.09
nuclear receptor coactivator 3	NM_024722	1.09
adenylyl cyclase-associated protein 2	AI761748	1.09
hypothetical protein MGC4707	NM_006366	1.09
KIAA0675 gene product	NM_024113	1.09
KIAA0738 gene product	BG502305	1.09
zinc finger protein 151 (pH2-87)	BC000609	1.09
cytochrome P450, subfamily IVF, polypeptide 11	AW574837	1.09
heat shock 70kD protein 6 (HSP70B)	NM_021187	1.09
KIAA0185 protein	X51757	1.09
SEX gene	AL547263	1.09
protein tyrosine phosphatase, receptor type, U	AB75453	1.09
Ras-induced senescence 1	U71075	1.09
endonuclease G-like 1	BF062629	1.09
	BF511976	1.09

hypothetical protein FLJ20170
 Consensus includes gb:BF155672 /FEA=EST /DB_XREF=gi:11050855 /DB_XREF=est:QV4-HT0894-290900-399-c04 /UG=Hs.146200 ESTs
 dystrophin myotonic-protein kinase
 killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 5
 Consensus includes gb:AV781453 /FEA=EST /DB_XREF=gi:10919301 /DB_XREF=est:AV781453 /CLONE=MOSBZA03 /UG=Hs.294014 ESTs
 egl-like module containing, mucin-like, hormone receptor-like sequence 1
 whigless-type MMTV integration site family, member 1
 myosin, heavy polypeptide 8, skeletal muscle, perinatal
 myosin binding protein C, slow type
 zinc finger protein 23 (K0X 16)
 thymine-DNA glycosylase
 hypothetical protein FLJ20635
 neuritin 3
 regenerating islet-derived 1 beta (pancreatic stone protein, pancreatic thread protein)
 xanthine dehydrogenase
 hypothetical protein FLJ10474
 ATPase, Ca++ transporting, plasma membrane 4
 hypothetical protein FLJ10508
 serologically defined colon cancer antigen 3
 son of sevenless (Drosophila) homolog 2
 TIA1 cytotoxic granule-associated RNA binding protein-like 1
 ttfelin 1
 KIA0814 gene product
 hypothetical protein PRO0813
 apolipoprotein (a) related gene C
 cyclin, basic protein of sperm head cytoskeleton 2
 tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase
 KIA0537 gene product
 hypothetical protein PRO2325
 proline-rich protein Bat1 subfamily 2
 Consensus includes gb:AA142942 /FEA=EST /DB_XREF=gi:1712320 /DB_XREF=est:z43c04.s1 /CLONE=IMAGE:504678 /UG=Hs.241507 ribosomal protein S6
 DiGeorge syndrome gene A
 arylsulfolase A
 RecQ protein-like 5
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 5
 KIA0565 gene product
 ELKL motif kinase
 solute carrier family 4, sodium bicarbonate cotransporter-like, member 10
 p53-induced protein
 Consensus includes gb:BG403760 /FEA=EST /DB_XREF=gi:13287238 /DB_XREF=est:602419627F1 /CLONE=IMAGE:4526599 /UG=Hs.158154 ESTs
 cytosolic acyl coenzyme A thioester hydrolase
 rad8 GTPase activating protein (GAP and centrosome-associated)
 transcription termination factor, mitochondrial
 CGI-29 protein
 ATP-binding cassette, sub-family F (GCN20), member 2
 Consensus includes gb:AU146809 /FEA=EST /DB_XREF=gi:11008330 /DB_XREF=est:AU146809 /CLONE=HEMBA1001564 /UG=Hs.287476 Homo sapiens cDNA FLJ12121 /AU146809
 hypothetical protein FLJ11342
 DKFZP434C171 protein
 tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase
 Consensus includes gb:AW969803 /FEA=EST /DB_XREF=gi:8159647 /DB_XREF=est:EST381881 /UG=Hs.105791 ESTs
 nuclear respiratory factor 1
 retinol binding protein 4, plasma
 Consensus includes gb:AU147800 /FEA=EST /DB_XREF=gi:11008321 /DB_XREF=est:AU147800 /CLONE=MAMMA1001745 /UG=Hs.296710 Homo sapiens cDNA FLJ12122 /AU147800
 hypothetical protein FLJ23588
 hypothetical protein FLJ20392
 Consensus includes gb:A1590063 /FEA=EST /DB_XREF=gi:4599101 /DB_XREF=est:t75d05.x1 /CLONE=IMAGE:2224137 /UG=Hs.235160 Homo sapiens mRNA, cDNA F.A1590063
 similar to KIA0010 gene product (H. sapiens)
 MAP/microtubule affinity-regulating kinase 1
 hypothetical protein FLJ11000
 Consensus includes gb:AL049710 /DEF=Human DNA sequence from clone 633H17 on chromosome 1p31.2-32.2. Contains a pseudogene similar to part of MTCO1 (Cytochrome AL049710
 Consensus includes gb:AW007532 /FEA=EST /DB_XREF=gi:5656310 /DB_XREF=est:ws52h07.x1 /CLONE=IMAGE:2500861 /UG=Hs.103391 Human insulin-like growth fr. AW007532
 hypothetical protein FLJ20550
 endothelial cell-specific molecule 1

carbonic anhydrase IV
 HMG-box containing protein 1
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3
 KIAA0894 gene product
 growth hormone secretagogue receptor
 protein tyrosine phosphatase, non-receptor type substrate 1
 melanoma cell adhesion molecule
 Consensus includes gb:AL136527 /DEF=Human DNA sequence from clone RP11-215B13 on chromosome 13 Contains the AKAP11 (A kinase (PRKA) anchor protein 11);
 Consensus includes gb:AV652328 /FEA=EST /DB_XREF=gi:3873342 /DB_XREF=est:AV652328 /CLONE=GLCCZC01 /UG=Hs.282488 ESTs
 calmodulin 1 (phosphorylase kinase, delta)
 hypothetical protein FLJ23403
 glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
 WW domain-containing oxidoreductase
 hypothetical protein FLJ11556
 Cluster Ind. N80935:207706.a1 /Homo sapiens cDNA, 3' end /clone=IMAGE:301402 /clone_end=3 /gb=N80935 /gi=1243636 /ug=Hs.22483 /len=527
 Consensus includes gb:AL138781 /DEF=Human DNA sequence from clone RP11-18H23 on chromosome 10. Contains the gene KIAA0204 (HSLX)
 corticotropin releasing hormone
 estrogen receptor 1
 major histocompatibility complex, class I, E
 geranylgeranyl diphosphate synthase 1
 lactase
 zinc finger protein 237
 lectin, galactoside-binding, soluble, 4 (galectin 4)
 Fc fragment of IgA, receptor for
 osteoglycin (osteoblast-inductive factor, mimecan)
 Consensus includes gb:AU145411 /FEA=EST /DB_XREF=gi:11006932 /DB_XREF=est:AU145411 /CLONE=HEMBA1004753 /UG=Hs.274546 /Homo sapiens cDNA FLJ10
 KIAA0869 protein
 transcription factor 3 (E2A Immunoglobulin enhancer binding factors E12/E47)
 galactose-1-phosphate uridylyltransferase
 hypothetical protein FLJ20802
 hypothetical protein MGC2383
 Consensus includes gb:AK023690.1 /DEF=Homo sapiens cDNA FLJ13628 fs, clone PLACE1011054 /FEA=mRNA /DB_XREF=gi:10435692 /UG=Hs.306656 /Homo sapiens
 hypothetical protein MGC3035
 patched homolog 2 (Drosophila)
 small inducible cytokine A5 (RANTES)
 KIAA1040 protein
 hypothetical protein FLJ12572
 Consensus includes gb:AK024890.1 /DEF=Homo sapiens cDNA: FLJ21237 fs, clone COL01114 /FEA=mRNA /DB_XREF=gi:10437303 /UG=Hs.308720 /Homo sapiens cD
 tumor necrosis factor receptor superfamily, member 18
 sterol O-acetyltransferase 2
 Fanconi anemia, complementation group F
 calcium channel, voltage-dependent, gamma subunit 1
 fibrinogen silencer binding protein
 Consensus includes gb:AK027246.1 /DEF=Homo sapiens cDNA: FLJ23593 fs, clone LNG14821 /FEA=mRNA /DB_XREF=gi:10440333 /UG=Hs.287749 /Homo sapiens cD
 hypothetical protein FLJ11850
 pregnancy specific beta-1-glycoprotein 1
 ribosomal protein L13
 Consensus includes gb:AL133571.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434F1135 (from clone DKFZp434F1135) /FEA=mRNA /DB_XREF=gi:5559148 /UG=Hs.11f
 acetylcholinesterase (YT blood group)
 hypothetical protein MGC2815
 T cell receptor gamma locus
 phosphatidylinositol transfer protein
 T cell receptor gamma constant 2
 cyclic AMP phosphoprotein, 19 kD
 hypothetical protein FLJ10241
 Huntingtin interacting protein E
 f-box and leucine-rich repeat protein 11
 telomerase-associated protein 1
 paired box gene 8
 camiline palmitoyltransferase II
 thrombospondin 1
 Interleukin 1, alpha
 GM2 ganglioside activator protein

heme binding protein 1
zinc finger protein, subfamily 1A, 5 (Pegasus)
Consensus includes gb:BC005286.1 /DEF=Homo sapiens, clone IMAGE3935812, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE3935812) /DB_Xt BC005286
chromosome 21 open reading frame 18
hypothetical protein FLJ22402
gb:AC276653.1 /DEF=Homo sapiens MAP1 light chain 3-like protein 2 mRNA, complete cds. /FEA=mRNA /PROD=MAP1 light chain 3-like protein 2 /DB_XREF=gqi:1362577 AF276659
Consensus includes gb:AL358589.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547C126 (from clone DKFZp547C126). /FEA=mRNA /DB_XREF=gqi:8655666 JUG=Hs.2638 AL358589
gamma-aminobutyric acid (GABA) B receptor, 1
Consensus includes gb:AL390143.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547N074 (from clone DKFZp547N074). /FEA=mRNA /DB_XREF=gqi:9368876 JUG=Hs.2842 AL390143
tripartite motif-containing 15
histone deacetylase 8
cell division cycle 25A
Consensus includes gb:AL137285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D2416 (from clone DKFZp434D2416). /FEA=mRNA /DB_XREF=gqi:6807740 JUG=Hs.30 AL137285
pregnancy-associated plasma protein A
occludin
protein phosphatase 5, catalytic subunit
hypothetical protein FLJ10040
period homolog 2 (Drosophila)
basic transcription factor 3
p53-regulated DDA3
Consensus includes gb:AL049646 /DEF=Human DNA sequence from clone RP4-568F9 on chromosome 20 Contains the ZNF133 (zinc finger protein 133 (clone pHZ-13)) /AL049646
DKFZP547G0910 protein
phosphatase and tetraol homolog (mutated in multiple advanced cancers 1)
zinc finger protein, Y-linked
Kallmann syndrome 1 sequence
glutamate-aminonitrile ligase (glutamine synthase)
keratin 18
aspartate beta-hydroxylase
thrombosane A2 receptor
RAB7, member RAS oncogene family
guanosine monophosphate reductase
plectin 1, intermediate filament binding protein, 500kD
transferrin
chitinase 1 (chitinotriase)
hypothetical protein MGC4365
GRB2-associated binding protein 2
protein kinase, AMP-activated, alpha 1 catalytic subunit
PHD finger protein 1
gb:AF130050.1 /DEF=Homo sapiens clone FLB3442 PRO0872 mRNA, complete cds. /FEA=mRNA /PROD=PRO0872 /DB_XREF=gqi:11493406 JUG=Hs.172154 Homo sap AF130050
acrosin
phosphatidylinositol-4-phosphate 5-kinase, type II, beta
KIAA1116 protein
KIAA0346 protein
testis transcript Y 9
KIAA0449 protein
Consensus includes gb:AU146999 /FEA=EST /DB_XREF=gqi:11008520 /DB_XREF=est:AU146999 /CLONE=HEMBB1002092 JUG=Hs.296688 Homo sapiens cDNA FLJ121
Consensus includes gb:AL577024 /FEA=EST /DB_XREF=gqi:12939748 /DB_XREF=est:AL577024 /CLONE=CS001082YN19 (3 prime) /JUG=Hs.7921 Homo sapiens mRNA; AL577024
DKFZP34K114 protein
canceroblastic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen)
keratin, hair, basic, 5
p53 regulated PA28 nuclear protein
KIAA1416 protein
ATP-binding cassette, sub-family A (ABC1), member 8
similar to hypothetical protein MNCb-2386
tumor endothelial marker 8
statin 1/oncogene protein 18
splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)
hypothetical protein FLJ20371
homologous to yeast nitrogen permease (candidate tumor suppressor)
calcium channel, voltage-dependent, alpha 1G subunit
hypothetical protein MGC13033
gb:AF212224.1 /DEF=Homo sapiens CLK4 mRNA, complete cds. /FEA=mRNA /PROD=CLK4 /DB_XREF=gqi:9437514 JUG=Hs.295231 Homo sapiens CLK4 mRNA, compk AF212224
tropomyosin 1 (alpha)
Z24727

stromal cell protein
 Consensus includes gb:8E613017 /FEA=EST /DB_XREF=gi:10245251 /DB_XREF=est:RC2-AN0062-140800-017-g03 /UG=Hs.124629 ESTs
 hypothetical protein FLJ20514
 Consensus includes gb:AK025194.1 /DEF=Homo sapiens cDNA: FLJ21541 flt, clone COL08166. /FEA=mRNA /DB_XREF=gi:1047857 /UG=Hs.308784 Homo sapiens cDNA: FLJ23356 flt
 retinoic acid repressible protein
 Consensus includes gb:AV711904 /FEA=EST /DB_XREF=gi:10731210 /DB_XREF=est:AV711904 /CLONE=DCAAIE08 /UG=Hs.277431 Homo sapiens cDNA: FLJ23356 flt
 Protein P3
 gb:BC004241.1 /DEF=Homo sapiens. Similar to laminin, alpha 4, clone MGC:10389. mRNA, complete cds. /FEA=mRNA /PROO=Similar to laminin, alpha 4 /DB_XREF=gi:BC004241
 KIAA0222 gene product
 AL110158
 U04845
 collagen, type IV, alpha 6
 U04845
 sarcoplasmic
 hypothetical protein FLJ21820
 quinine oxidoreductase homolog
 PAS-serine/threonine kinase
 trefoil factor 2 (spasmolytic protein 1)
 KIAA0878 protein
 ELL-RELATED RNA POLYMERASE II, ELONGATION FACTOR
 host cell factor homolog
 BCL2-interacting killer (apoptosis-inducing)
 ISL1 transcription factor, LIM/homeodomain, (Islet-1)
 signal-regulatory protein beta 1
 Consensus includes gb:AF217536.1 /DEF=Homo sapiens truncated mevalonate kinase mRNA, partial cds, alternatively spliced. /FEA=mRNA /PROO=truncated mevalonate
 GS3955 protein
 myotubularin related protein 2
 syndecan 4 (amphiglycan, ryudocan)
 arginase, type II
 neurotrophic tyrosine kinase, receptor, type 2
 actin cross-linking factor
 PRO1575 protein
 Consensus includes gb:AW451236 /FEA=EST /DB_XREF=gi:6992012 /DB_XREF=est:UI-H-B13-ah-a-07-0-UI.s1 /CLONE=IMAGE:2736661 /UG=Hs.15535 Homo sapiens
 hypothetical protein FKSG44
 protocadherin gamma subfamily A, 3
 alkaline phosphatase, placental (Regan isozyme)
 phosphatidylinositol glycan, class F
 solute carrier family 19 (folate transporter), member 1
 UHS KerB
 v-K-ras2 Kirsten rat sarcoma 2 viral oncogene homolog
 hypothetical protein FLJ20059
 fragile X mental retardation, autosomal homolog 2
 Consensus includes gb:AW972855 /FEA=EST /DB_XREF=gi:8162701 /DB_XREF=est:EST384950 /UG=Hs.292853 ESTs
 phosphate cytidylyltransferase 1, choline, alpha isoform
 cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 13
 c-myc promoter-binding protein
 TAR (HIV) RNA binding protein 1
 hypothetical protein FLJ23407
 thrombospondin 1
 forkhead box M1
 fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson
 Consensus includes gb:AV194543 /FEA=EST /DB_XREF=gi:6473381 /DB_XREF=est:tb27c02.s1 /CLONE=IMAGE:2577506 /UG=Hs.234573 Homo sapiens mRNA for TL
 hypothetical protein FLJ22623
 zinc finger protein 35 (clone HF.10)
 KIAA0663 gene product
 KIAA0752 protein
 Cluster Ind. N36928;yy08a06.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:273538 /clone_end=3 /gb=N36928 /gi=1158070 /ug=Hs.33540 /len=562
 KIAA0356 gene product
 Consensus includes gb:AF007147.1 /DEF=Homo sapiens clone 23712 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852625 /UG=Hs.113166 Homo sapiens clone 23712 AF007147
 HNK-1 sulfotransferase
 cholinergic receptor, nicotinic, alpha polypeptide 10
 myotubular myopathy 1
 polymerase I and transcript release factor
 interleukin 11

gb:AL136548.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G18121 (from clone DKFZp761G18121); complete cds. /FEA=mRNA /GEN=DKFZp761G18121 /PROD=hyp
 pancreatic elastase 1B
 1.08 NM_015849
 hypothetical protein FLJ20070
 1.08 NM_017652
 Consensus includes gb:U80773.1 /DEF=Human EST clone 42944 mariner transposon Hsmar1 sequence /FEA=mRNA /DB_XREF=gi:2231376 /UG=Hs.121580 Human ES
 1.08 BG111661
 goji autoantigen, gojiin subfamily a, 1
 1.08 NM_008808
 gamma-aminobutyric acid (GABA) A receptor, alpha 3
 1.08 NM_014237
 a disintegrin and metalloproteinase domain 18
 1.08 AWT48405
 neuritin 1
 1.08 NM_030808
 LIS1-interacting protein NUDEL; endonucleopeptidase A
 1.08 NM_007350
 pleckstrin homology-like domain, family A, member 1
 1.08 BC001090
 KIAA1668 protein
 1.08 AB014607
 KIAA0707 protein
 1.08 NM_016265
 G10T-3 for gonadotropin inducible transcription repressor-3
 1.08 AG02100
 dedual protein induced by progesterone
 1.08 AF068266
 core-binding factor, runt domain, alpha subunit 2; translocated to, 2
 1.08 NM_006606
 retinoblastoma binding protein 9
 1.08 AW007137
 hypothetical protein FLJ10342
 1.08 NM_003186
 transgelin
 1.08 NM_004726
 RALBP1 associated Eps domain containing 2
 1.08 NM_024300
 hypothetical protein MGC2217
 1.08 NM_001297
 cyclic nucleotide gated channel beta 1
 1.08 NM_002114
 human immunodeficiency virus type 1 enhancer binding protein 1
 1.08 AA911235
 macrophage stimulating, pseudogene 9
 1.08 NM_001255
 CDC20 cell division cycle 20 homolog (S. cerevisiae)
 1.08 AJ133788
 LIM domain binding 3
 1.08 NM_025257
 NG22 protein
 1.08 AA723370
 CGL-105 protein
 1.08 A073549
 estrogen receptor 1
 1.08 Z70519
 tumor necrosis factor receptor superfamily, member 6
 1.08 NM_024759
 hypothetical protein FLJ13955
 1.08 NM_0110164
 Consensus includes gb:AL110164.1 /DEF=Homo sapiens mRNA; cDNA DKFZp58610324 (from clone DKFZp58610324). /FEA=mRNA /DB_XREF=gi:5817069 /UG=Hs.1937
 1.08 AL110164
 hypothetical protein PRO2012
 1.08 AA422049
 hypothetical protein DKFZp547M138 similar to widely-interspaced zinc finger motifs
 1.08 AK022326
 Consensus includes gb:AK022326.1 /DEF=Homo sapiens cDNA FLJ12264 fls, clone MAMMA1001600. /FEA=mRNA /DB_XREF=gi:10433699 /UG=Hs.288481 Homo sapi
 1.08 AF349719
 trophinin
 1.08 NM_001390
 dystrobrevin, alpha
 1.08 BF444943
 ubiquitin specific protease 24
 1.08 L13605
 cholecystokinin A receptor
 1.08 AF133425
 tetraspan 1
 1.08 AL041852
 KIAA1464 protein
 1.08 NM_003322
 bubby like protein 1
 1.08 NM_000165
 serine (or cysteine) proteinase inhibitor, clade D (heparin cofactor), member 1
 1.08 AL139112
 Consensus includes gb:AL139112 /DEF=Human DNA sequence from clone GS1-103B18 on chromosome Xq27.1-27.3 Contains ESTs, STSs and GSSs. Contains a novel
 1.08 U14383
 mucin 8, tracheobronchial
 1.08 AB016900
 HGC8.1.1 protein
 1.08 NM_000711
 bone gamma-carboxyglutamate (gla) protein (osteocalcin)
 1.07 AF001690
 exostosins (multiple)-like 3
 1.07 AL045800
 trinuclidic repeat containing 15
 1.07 NM_025102
 hypothetical protein FLJ21487
 1.07 AF106934
 thyroid hormone receptor-associated protein, 95-kD subunit
 1.07 NM_020387
 CATX-8 protein
 1.07 AA633841
 surfactant, pulmonary-associated protein C
 1.07 AL136823
 thyroid hormone receptor coactivating protein
 1.07 NM_003297
 nuclear receptor subfamily 2, group C, member 1
 1.07 NM_024028
 hypothetical protein MGC3265
 1.07 M13452
 lamin A/C
 1.07 NM_026894
 Consensus includes gb:AK026894.1 /DEF=Homo sapiens cDNA: FLJ23041 fls, clone LNG002288. /FEA=mRNA /DB_XREF=gi:10439604 /UG=Hs.306865 Homo sapiens cC
 1.07 NM_001488
 glutamate-cysteine ligase, catalytic subunit
 1.07 BC000076
 cyclin D1 (PRAD1; parathyroid adenomatosis 1)
 1.07 AK021487
 Consensus includes gb:AK021487.1 /DEF=Homo sapiens cDNA FLJ11425 fls, clone HEMBA1001051. /FEA=mRNA /DB_XREF=gi:10432680 /UG=Hs.306597 Homo sapi
 1.07 BC005253
 Ketch motif containing protein
 1.07 AK026040
 KIAA0876 protein
 1.07 A1743713
 Notch homolog 4 (Drosophila)

casepase 9, apoptosis-related cysteine protease	U60521	1.07
tripartite motif-containing 15	U34249	1.07
transforming growth factor, beta receptor I (activin A receptor type II-like kinase, 53kD)	NM_004612	1.07
decdinase, dectinomyone, type III	NM_001382	1.07
succinate-CoA ligase, GDP-forming, beta subunit	AV650470	1.07
ATP-binding cassette, sub-family D (ALD), member 4	NM_005050	1.07
serine/threonine kinase 17a (apoptosis-inducing)	AW183478	1.07
hypothetical protein FLJ14360	BG325646	1.07
myosin, light polypeptide 5, regulatory	L03785	1.07
contactin associated protein-like 2	AU144598	1.07
PAB23, member PAS oncogene family	NM_016277	1.07
estrogen regulated gene 1	NM_022034	1.07
cadherin 6, type 2, K-cadherin (fetal kidney)	AU151463	1.07
catenin, beta interacting protein 1	NM_020248	1.07
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9	AF248968	1.07
neuronal pentraxin receptor	AL162057	1.07
XO4603 B subunit thrc, thrb genes corresponding to nucleolides 248-2228 of XO4603 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	XO4603	1.07
Consensus includes gb:BF223370 /FEA=EST /DB_XREF=gi:1130547 /DB_XREF=est:q87603.x1 /CLONE=IMAGE:3705316 /UG=Hs.90449 Human clone 23908 mRNA	BF223370	1.07
phosphatidylserine-specific phospholipase A1 alpha	NM_015900	1.07
excision repair cross-complementing rodent repair deficiency, complementation group 6	NM_000124	1.07
insulin-like 4 (placenta)	NM_002195	1.07
Consensus includes gb:U79249.1 /DEF=Human clone 23839 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710196 /UG=Hs.76362 Human clone 23839 mRNA sequence	U79249	1.07
ribosomal protein S6 kinase, 80kD, polypeptide 6	AF184965	1.07
chordin-like	AL049176	1.07
protocadherin gamma subfamily C, 3	NM_002588	1.07
transferrin receptor (p90, CD71)	BC001188	1.07
tumor protein p53 binding protein, 1	NM_005657	1.07
metallothionein 3 (growth inhibitory factor (neurotrophic))	NM_005954	1.07
interleukin 7	NM_000880	1.07
pyrimidinergic receptor P2Y, G-protein coupled, 4	NM_002585	1.07
hypothetical protein	NM_016101	1.07
lymphocyte antigen 8 complex, locus D	NM_003695	1.07
vav 3 oncogene	AF118887	1.07
leukocyte-associated Ig-like receptor 2	NM_002288	1.07
KIA0381 protein	BF513244	1.07
KIA0197 protein	D83781	1.07
hypothetical protein NGC2963	NM_031298	1.07
KIAA1718 protein	BE217882	1.07
serine/threonine kinase 38 like	AW779556	1.07
B-cell CLL/lymphoma 7A	BC002829	1.07
Integrin, beta 4	AA808063	1.07
exostosins (multiple)-like 2	AF000416	1.07
ubiquitin specific protease 21	NM_012475	1.07
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila), translocated to, 3	AV756536	1.07
kinase-like 7	NM_020242	1.07
Consensus includes gb:AU145711 /FEA=EST /DB_XREF=gi:11007232 /DB_XREF=est:AU145711 /CLONE=HEMBA1005588 /UG=Hs.301008 Homo sapiens cDNA FLJ11	AU145711	1.07
melanoma antigen, family A, 12	BC003408	1.07
plihn, desmosome associated protein	BF508848	1.07
transitional epithelia response protein	NM_013319	1.07
PTH-responsive osteosarcoma B1 protein	U87408	1.07
Interleukin 6 signal transducer (gp130, oncostatin M receptor)	BE856546	1.07
DKF ZP434N014 protein	AA813332	1.07
single-strand selective monofunctional uracil DNA glycosylase	NM_014311	1.07
C-terminal binding protein 1	BF337195	1.07
myosin, heavy polypeptide 2, skeletal muscle, adult	NM_017534	1.07
STO protein	NM_014070	1.07
casepase 10, apoptosis-related cysteine protease	U88214	1.07
Consensus includes gb:AL157375 /DEF=Human DNA sequence from clone RP1-133P16 on chromosome 6 Contains the ARF1 (ADP-ribosylation factor 1) gene, ESTs, ST, AL157375	AL157375	1.07
hypothetical protein FLJ21472	NM_024837	1.07
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls, clone MAMMA1001858, /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie	AK022363	1.07
dihydropyrate dehydrogenase	BE789211	1.07
hypothetical protein	NM_016399	1.07
KIAA0707 protein	AK023537	1.07

beta-1,4 mannosyltransferase	1.07
Consensus includes gb:AF103295.1 /DEF=Homo sapiens clone N97 immunoglobulin heavy chain variable region mRNA, partial cds. /FEA=mrna /PROD=immunoglobulin	1.07
nifunlin 1	1.07
AE binding protein 1	1.07
KIAA0982 protein	1.07
complement component 3	1.07
tumor necrosis factor (ligand) superfamily, member 6	1.07
hypothetical protein FLJ12377	1.07
phosphatidylinositol glycan, class L	1.07
G protein-coupled receptor 25	1.07
hypothetical protein FLJ11743	1.07
Consensus includes gb:AA975427 /FEA=EST /DB_XREF=gi:3151219 /DB_XREF=estcoq28902.s1 /CLONE=IMAGE:1587698 /UG=Hs.128434 Homo sapiens ELISC-1 mRNA	1.07
solute carrier family 2 (facilitated glucose transporter), member 3	1.07
calcium channel, voltage-dependent, beta 1 subunit	1.07
aminoacyl-tRNA synthetase	1.07
small inducible cytokine subfamily B (Cys-X-Cys), member 10	1.07
CGI-44 protein, sulfide dehydrogenase like (yeast)	1.07
catenin (cadherin-associated protein), alpha-like 1	1.07
nuclear receptor subfamily 5, group A, member 2	1.07
golg-specific brefeldin A resistance factor 1	1.07
caspace 10, apoptosis-related cysteine protease	1.07
KIAA0202 protein	1.07
PTG-binding protein	1.07
myosin, heavy polypeptide 11, smooth muscle	1.07
protocadherin gamma subfamily A, 9	1.07
protein kinase C binding protein 1	1.07
carbohydrate (keratin sulfate Gal-6) sulfotransferase 1	1.07
cholinergic receptor, nicotinic, delta polypeptide	1.07
superficial viral-like activity 2-like (S. cerevisiae)	1.07
deleted in lung and esophageal cancer 1	1.07
breakpoint cluster region	1.07
vesicovacuolar intestinal peptide receptor 2	1.07
phosphoinositide-specific phospholipase C-beta 1	1.07
enhancer of rudimentary homolog (Drosophila)	1.07
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson	1.07
polyubiquitin-like protein	1.07
seven transmembrane protein TM7SF3	1.07
uncharacterized hematopoietic stem/progenitor cells protein MDS028	1.07
nucleosome assembly protein 1-like 1	1.07
glutamate receptor, ionotropic, N-methyl D-aspartate 1	1.07
spinocerebellar ataxia 8	1.07
terranectin (plasminogen binding protein)	1.07
disrupted in bipolar disorder 1	1.07
deleted in cancer 1: RNA helicase HDB/DICE1	1.07
KIAA0274 gene product	1.07
UDP glycosyltransferase 2 family, polypeptide A1	1.07
ring finger protein 15	1.07
G protein-coupled receptor kinase 2-like (Drosophila)	1.07
KIAA0210 gene product	1.07
hypothetical protein FLJ11827	1.07
Consensus includes gb:AP000693 /DEF=Homo sapiens genomic DNA, chromosome 21q22.2, PCR fragment from BAC clone XBT79C11, CBR1-HLCS region /FEA=mrna /PROD=transcript	1.07
interferon gamma receptor, binding protein 2 (lprin beta 2)	1.07
KIAA1117 protein	1.07
KATA box binding protein (TBP)-associated factor, RNA polymerase I, A, 46kD	1.07
Golg apparatus protein 1	1.07
ecotropic viral integration site 1	1.07
PMS2 postmeiotic segregation increased 2 (S. cerevisiae)	1.07
keratin, hair, acidic, 1	1.07
calumenin	1.07
retinol X receptor, beta	1.07
CGI-48 protein	1.07
mannose-binding lectin (protein C) 2, soluble (opsonic defect)	1.07
polymerase (RNA) mitochondrial (DNA directed)	1.07
BF057617	1.07

synaptrophin, alpha 1(dystrophin-associated protein A1, 59kD, acidic component)
 epidermal growth factor receptor pathway substrate 8
 hypothetical protein FLJ12078
 hypothetical protein FLJ12078
 thoredoxin domain-containing
 integrin, beta 3 (platelet glycoprotein IIa, antigen CD61)
 hypothetical protein FLJ21308
 Consensus includes gb:AL390137.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D179 (from clone DKFZp434D179). /FEA=mRNA /DB_XREF=gi:9368847 /UG=Hs.3065
 belaine-homocysteine methyltransferase 2
 purinergic receptor P2X, ligand-gated ion channel, 1
 LIM homeobox protein 8
 homeodomain interacting protein kinase 3
 cardioembryonic antigen-related cell adhesion molecule 7
 nucleoporin 88KD
 contactin 5
 p53-induced protein PIGPC1
 activin A receptor type II-like 1
 UDP-glucose ceramide glucosyltransferase-like 2
 KIAA1503 protein
 aquaporin 5
 claudin 16
 PHD finger protein 1
 Consensus includes gb:AL137713.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434I0523 (from clone DKFZp434I0523). /FEA=mRNA /DB_XREF=gi:6808127 /UG=Hs.3064 /AL137713
 amyloid beta (A4) precursor protein-binding, family A, member 2 (X11-like)
 gb:NM_003902.1 /DEF=Homo sapiens far upstream element (FUSE) binding protein 1 (FUBP1), mRNA. /FEA=mRNA /GEN=FUBP1 /PROD=far upstream element-binding
 BC000585
 solute carrier family 21 (organic anion transporter), member 11
 R3H domain (binds single-stranded nucleic acids) containing
 plastin 3 (T isoform)
 apolipoprotein L, 3
 adrenal gland protein AD-004
 KIAA0096 protein
 A1Pase, Cx++ transporting, plasma membrane 3
 activating transcription factor 5
 PTH-responsive osteosarcoma B1 protein
 popeye protein 3
 novel RGD-containing protein
 hypothetical protein FLJ13852
 RAB2, member RAS oncogene family
 Consensus includes gb:A1769888 /FEA=EST /DB_XREF=gi:5236197 /DB_XREF=est:W2511.x1 /CLONE=IMAGE:2403885 /UG=Hs.180737 Hmo sapiens clone 23664 an
 A1769888
 hypothetical protein FLJ11585
 KIAA0232 gene product
 DnaJ (Hsp40) homolog, subfamily C, member 8
 BNS1 (BHK21) temperature sensitivity complementing
 solute carrier family 31 (copper transporters), member 1
 reversion-inducing-cysteine-rich protein with kazal motifs
 KIAA0805 protein
 M245378 subtilisin phageA genes corresponding to nucleolides 2017-3334 of M24537 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
 frizzled homolog 8 (Drosophila)
 Consensus includes gb:A1471375 /FEA=EST /DB_XREF=gi:4333465 /DB_XREF=est:tm10c12.x1 /CLONE=IMAGE:2156182 /UG=Hs.80206 glucose-6-phosphate dehydrog
 L38969
 thrombospondin 3
 short stature homeobox 2
 mitochondrial ribosomal protein S26
 Consensus includes gb:AL137378.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K1128 (from clone DKFZp434K1128). /FEA=mRNA /DB_XREF=gi:6807908 /UG=Hs.30
 AL121905
 CD38 antigen (collagen type I receptor, thrombospondin receptor)-like 1
 splicing factor 3b, subunit 4, 49KD
 putative translation initiation factor
 villin-like
 S100 calcium binding protein A7 (psoriasin 1)
 procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3
 hypothetical protein FLJ11078
 apolipoprotein E
 cyclin T2
 tyrosine hydroxylase
 UDP-N-acetyl-alpha-D-galactosamine:(N-acetylneuraminyl)-galactosylglucosylceramide N-acetylgalactosaminyltransferase (GalNAc-T)
 ALD50358

Rho guanine exchange factor (GEF) 11
 Consensus includes gb:AK024688.1 /DEF=Homo sapiens cDNA: FLJ12125 fls, clone COL00526, /FEA=mrRNA /DB_XREF=gi:10437276 /UG=Hs.306711 Homo sapiens cDNA: AK024688
 carboxypeptidase E
 Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA: FLJ12241 fls, clone MAMMA1001274, /FEA=mrRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens cDNA: AK022303
 erythroid differentiation and denudation factor 1
 DNFXP34A0131 protein
 WNT1 inducible signaling pathway protein 1
 deleted in malignant brain tumors 1
 isochlorate dehydrogenase 3 (NAD+) alpha
 GAS2-related on chromosome 22
 adenylylase cyclase 2 (brain)
 KIAA1484 protein
 caspase recruitment domain protein 14
 hypothetical protein similar to preferentially expressed antigen of melanoma
 nuclear receptor co-repressor 1
 ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome
 Consensus includes gb:AU148184 /FEA=EST /DB_XREF=gi:11009705 /DB_XREF=est:AU148184 /CLONE=MAMMA1002820 /UG=Hs.287529 Homo sapiens cDNA: FLJ12 AU148184
 CS box-containing WD protein
 RAN binding protein 2-like 1
 H1 histone family, member 3
 Consensus includes gb:AU073983 /FEA=EST /DB_XREF=gi:3400627 /DB_XREF=IMAGE:1670790 /UG=Hs.298553 Homo sapiens, clone IMAGE: AU073983
 fatty acid desaturase 1
 cAMP responsive element modulator
 syntaxin binding protein 3
 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta
 PTH-responsive osteosarcoma B1 protein
 paired basic amino acid cleaving enzyme (furin, membrane associated receptor protein)
 hypothetical protein FLJ10955
 dopamine receptor D5
 pyruvate dehydrogenase kinase, isoenzyme 4
 apolipoprotein A-II
 Consensus includes gb:AL049244.1 /DEF=Homo sapiens mRNA: cDNA: DKFZp564C163 (from clone DKFZp564C163), /FEA=mrRNA /DB_XREF=gi:499979 /UG=Hs.3020 AL049244
 Consensus includes gb:AU157200 /FEA=EST /DB_XREF=gi:11018721 /DB_XREF=est:AU157200 /CLONE=PLACE1006617 /UG=Hs.301390 Homo sapiens cDNA: FLJ135 AU157200
 neurofibromin 2 (bilateral acoustic neuroma)
 neural poly(pyrimidine) tract binding protein
 galactose-4-epimerase, UDP-
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 1
 collagen, type IV, alpha 6
 killer cell lectin-like receptor subfamily C, member 2
 tumor necrosis factor receptor superfamily, member 9
 hypothetical protein FLJ10378
 adrenergic, beta, receptor kinase 1
 hypothetical protein FLJ10252
 cystatin SA
 solute carrier family 6 (neurotransmitter transporter, dopamine), member 3
 KIAA1077 protein
 hypothetical protein FLJ10339
 egrin
 naturalistic peptide receptor C/guanylate cyclase C (atrionatriuretic peptide receptor C)
 cytoline receptor-like factor 2
 hypothetical protein FLJ20531
 Consensus includes gb:T53900 /FEA=EST /DB_XREF=gi:655761 /DB_XREF=est:y83e08.r1 /CLONE=IMAGE:77798 /UG=Hs.172035 hypothetical protein similar to mouse: T53900
 Consensus includes gb:AI382123 /FEA=EST /DB_XREF=gi:4194904 /DB_XREF=est:y83e09.x1 /CLONE=IMAGE:2087416 /UG=Hs.182919 ESTs, Highly similar to AS925: AI382123
 Rho guanine nucleotide exchange factor (GEF) 10
 zinc finger protein 225
 hypothetical protein MGC9084
 cholinergic receptor, nicotinic, alpha polypeptide 3
 chromodomain helicase DNA binding protein 1
 netrin G1
 NDRG family, member 4
 5-methyltetrahydrofolate-homocysteine methyltransferase reductase
 immature colon carcinoma transcript 1
 retinol dehydrogenase 5 (11-cis and 9-cis)

chromosome 1 open reading frame 16
 G protein-coupled receptor 51
 Consensus includes gb:AU147084 /FEA=EST /DB_XREF=gi:11008605 /DB_XREF=est:AU147084 /CLONE=HEMBB1002371 /UG=Hs.287482 Homo sapiens cDNA FLJ121
 cameline palmitoyltransferase II
 protein tyrosine phosphatase, non-receptor type 21
 Ca²⁺-dependent activator protein for secretion
 cd. 1A
 hemoglobin, theta 1
 serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor), member 1
 involucrin
 ADP-ribosyltransferase 1
 similar to RhoGAP
 DKFZP434C212 protein
 growth arrest-specific 1
 CGI-14 protein
 Consensus includes gb:AL359583.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D0328 (from clone DKFZp434D0328) /FEA=mRNA /DB_XREF=gi:8655619 /UG=Hs.30
 immunoglobulin lambda joining 3
 suppressor of Ty 4 homolog 1 (S. cerevisiae)
 protocadherin beta 17 pseudogene
 Rhesus blood group, CcEe antigens
 Consensus includes gb:AL388866 /DEF=Human DNA sequence from clone RP1-237C24 on chromosome 20 Contains the 3 end of the SLC23A1 gene encoding solute car
 heparan sulfate (glucosamine) 3-O-sulfotransferase 2
 ADP-ribosylation factor-like 7
 KIAA0848 protein
 RAB31, member RAS oncogene family
 ring finger protein 11
 insulin-like growth factor binding protein 1
 insulin-like growth factor binding protein 1
 T cell receptor alpha locus
 RAB5C, member RAS oncogene family
 cyclokeratin type II
 SRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal)
 solute carrier family 28 (sodium-coupled nucleoside transporter), member 2
 hypothetical protein FLJ10143
 TANK-binding kinase 1
 KIAA0141 gene product
 like mouse brain protein E46
 putative G-protein coupled receptor
 potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2
 Consensus includes gb:AU144324 /FEA=EST /DB_XREF=gi:11005845 /DB_XREF=est:AU144324 /CLONE=HEMBA1001570 /UG=Hs.306599 Homo sapiens cDNA FLJ11-
 matrix metalloproteinase 19
 Cluster incl. A1W033091:wr03d04.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2480455 /clone_end=3 /gb=AW003091 /gi=5849929 /ug=Hs.6217 /len=813
 EGF-like domain, multiple 5
 hypothetical gene CQ018
 retinol binding protein 1, cellular
 Consensus includes gb:AU153979 /FEA=EST /DB_XREF=gi:1015500 /DB_XREF=est:AU153979 /CLONE=NT2RP3004470 /UG=Hs.293928 Homo sapiens cDNA FLJ142
 zinc finger protein 235
 thyroid hormone receptor, beta (erythroblastic leukemia viral (v-erb-a) oncogene homolog 2, avian)
 KIAA0649 gene product
 Mouse Mammary Tumor Virus Receptor homolog
 deoxyribonuclease II-ka 2
 KIAA0383 protein
 KIAA0472 protein
 heterogeneous nuclear ribonucleoprotein H2 (H')
 heparan sulfate 2-O-sulfotransferase
 collagen, type V, alpha 1
 transmembrane 4 superfamily member (tetraspan NET-7)
 cadherin 4, type 1, R-cadherin (retinal)
 adrenergic, alpha-1D, receptor
 syndecan 3 (N-syndecan)
 KIAA0379 protein
 KIAA0335 gene product
 islet amyloid polypeptide

coagulation factor II (thrombin) receptor-like 3	NM_003950	1.06
collagen, type IV, alpha 3 (Goodpasture antigen)	M81379	1.06
KIAA0500 protein	AW028110	1.06
KIAA0523 protein	AL13046	1.06
KIAA0328 protein	AW003635	1.06
hypothetical protein PRO2188	NM_018621	1.06
Consensus includes gb:AW026481 /FEA=EST /DB_XREF=gi:5860011 /DB_XREF=est:ww14a04.x1 /CLONE=IMAGE:2529486 /UG=Hs.48396 ESTs, Moderately similar to	AW026481	1.06
regulator of G-protein signalling 4	NM_005613	1.06
SCN Circadian Oscillatory Protein (SCOP)	AB011178	1.06
short stature homeobox	NM_000451	1.06
aquaporin 3	AB001325	1.06
tripartite motif-containing 15	U34249	1.06
hypothetical protein PRO1598	NM_018503	1.06
caspase 8, apoptosis-related cysteine protease	NM_001228	1.06
gliome-amplified sequence-41	NM_006530	1.06
chloride channel 3	AA802571	1.06
chromosome 6 open reading frame 35	NM_018452	1.06
G protein-coupled receptor 24	NM_005297	1.06
hypothetical protein FLJ20116	AA469071	1.06
gb:AF130053.1 /DEF=Homo sapiens clone FLB4228 PRO1095 mRNA, complete cds. /FEA=mRNA /PROD=PRO1095 /DB_XREF=gi:11493412 /UG=Hs.302145 Homo sap	AF130053	1.06
teratocapitula repeat domain 3	NM_003316	1.06
tumor protein p53 binding protein, 2	NM_005426	1.06
caspase 7, apoptosis-related cysteine protease	NM_001227	1.06
jerky homolog (mouse)	AF072466	1.06
homeo box D3	NM_006898	1.06
KIAA0229 protein	AI890523	1.06
potassium voltage-gated channel, KOT-like subfamily, member 1	NM_000218	1.06
Consensus includes gb:N31716 /FEA=EST /DB_XREF=gi:1152115 /DB_XREF=est:yl5c12.s1 /CLONE=IMAGE:271318 /UG=Hs.31754 ESTs	N31716	1.06
Consensus includes gb:AU118026 /FEA=EST /DB_XREF=gi:109330043 /DB_XREF=est:AU118026 CLONE=HEMBA1002729 /UG=Hs.289068 Homo sapiens cDNA FLJ111	AU118026	1.06
Consensus includes gb:AW300309 /FEA=EST /DB_XREF=gi:5709909 /DB_XREF=est:xx59608.x1 /CLONE=IMAGE:2773958 /UG=Hs.27453 ESTs, Moderately similar to G	AW300309	1.06
carbonic anhydrase IX	NM_001216	1.06
potassium inwardly-rectifying channel, subfamily J, member 5	D50134	1.06
hypothetical protein MGC2869	BC005856	1.06
protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte)	NM_002830	1.06
melanoma antigen, family B, 2	AA058636	1.06
rodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4, autosomal dominant)	NM_016526	1.06
GS15	R25692	1.06
Consensus includes gb:R25692 /FEA=EST /DB_XREF=gi:781827 /DB_XREF=est:yg44112.2 /CLONE=IMAGE:35277 /UG=Hs.224829 ESTs	AB032983	1.06
KIAA1137 protein	AI338653	1.06
G protein-coupled receptor kinase 5	NM_002364	1.06
NEL-like 1 (chicken)	NM_006157	1.06
kalikrein 2, prostatic	AA595465	1.06
testis enhanced gene transcript (BAX inhibitor 1)	NM_003217	1.06
related RAS viral (r-ras) oncogene homolog 2	AI753792	1.06
glypican 1	AI354864	1.06
hypothetical protein FLJ13491	NM_024623	1.06
hypothetical protein FLJ21616	NM_024587	1.06
lysyl oxidase	NM_002317	1.06
aldehyde dehydrogenase 8 family, member A1	NM_022568	1.06
hypothetical protein FLJ21313	AW575493	1.06
interactin 1 (SH3 domain protein)	AF114486	1.06
core-binding factor, runt domain, alpha subunit 2; translocated to, 2	NM_005093	1.06
porcupine	NM_022825	1.06
secretory leukocyte protease inhibitor (antileukoprotease)	NM_003064	1.06
fertilization 1 homolog a	AB62658	1.06
Consensus includes gb:BE857715 /FEA=EST /DB_XREF=gi:10372018 /DB_XREF=est:7g46g03.x1 /CLONE=IMAGE:3309556 /UG=Hs.8982 ESTs, Highly similar to KIAA11	BE857715	1.06
DNFZP564M14.16 protein	BF981643	1.06
nuclear factor related to kappa B binding protein	AI887378	1.06
SRY (sex determining region Y)-box 30	NM_007017	1.06
PR domain containing 2, with ZNF domain	NM_012231	1.06
nucleoporin 214KD (CAIN)	D14889	1.06
avastases (multiple) 2	AA186245	1.06
sperm associated antigen 6	AF078363	1.06

antigen identified by monoclonal antibodies 12E7, F21 and O13
Consensus includes gb:A1733515 /FEA=EST /DB_XREF=gi:5054676 /DB_XREF=est:ou99e05.x5 /CLONE=IMAGE:1635988 /UG=Hs.148907 Homo sapiens mRNA; cDNA
pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1)
kynureninase (L-tryptophan hydrolase)
peroxiredoxin 3
ubiquitin specific protease 14 (iRNA-guanine transglycosylase)
H4 histone family, member M
CED-6 protein
phosphoinositide-3-kinase, class 2, beta polypeptide
cytochrome c oxidase subunit VIa polypeptide 2
hypothetical protein DKFZp564L2423
alpha-methylacyl-CoA racemase
cyclin-dependent kinase 2
NADPH oxidase 1
TRF2-interacting telomeric RAP1 protein
RAS guanyl releasing protein 2 (calcium and DAG-regulated)
Wolf-Hirschhorn syndrome candidate 2
KIAA0144 gene product
book2 protein
chimerin (chimerin 1)
Consensus includes gb:AJ235383.1 /DEF=Homo sapiens mRNA for immunoglobulin heavy chain variable region, ID 31. /FEA=mRNA /GEN=IGHV /PROD=immunoglobulin
Consensus includes gb:A1357616 /FEA=EST /DB_XREF=gi:4109237 /DB_XREF=est:qu20g08.x1 /CLONE=IMAGE:1985374 /UG=Hs.101651 Homo sapiens mRNA; cDNA
collagen, type V, alpha 3
neural precursor cell expressed, developmentally down-regulated 4-like
retinitis pigmentosa GTPase regulator
nucleosomal binding protein 1
myosin IXb
glutamate receptor, metabotropic 2
elafin
serine/arginine repetitive matrix 1
RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)
cytochrome P450, subfamily I1C (mephenytoin 4-hydroxylase), polypeptide 18
KIAA0471 gene product
Consensus includes gb:A1561253 /FEA=EST /DB_XREF=gi:4511584 /DB_XREF=est:q27g06.x1 /CLONE=IMAGE:2210074 /UG=Hs.306649 Homo sapiens cDNA FLJ1340
hypothetical protein FLJ12988
5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)
KIAA0892 protein
serologically defined breast cancer antigen NY-BR-20
hypothetical protein MGC10798
cAMP responsive element binding protein 1
Consensus includes gb:AL036840 /FEA=EST /DB_XREF=gi:5408335 /DB_XREF=est:DKFZp564N1963_s1 /CLONE=DKFZp564N1963 /UG=Hs.22370 Homo sapiens mRNA
syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)
ribosomal protein L18
ras responsive element binding protein 1
hypothetical protein FLJ20654
ubiquitin specific protease 18
parvin, beta
lung type-I cell membrane-associated glycoprotein
hypothetical protein FLJ20584
etoposide-induced mRNA
ubiquitously transcribed tetranucleotide repeat gene, X chromosome
hemochromatosis
KIAA1037 protein
GAS2-related on chromosome 22
Consensus includes gb:A1875944 /FEA=EST /DB_XREF=gi:4650775 /DB_XREF=est:y65d07.x1 /CLONE=IMAGE:2283949 /UG=Hs.295963 ESTs
lipocalin 1 (protein migrating faster than albumin, tear prealbumin)
Toll-like receptor 8
cyclin-dependent kinase (CDC2-like) 10
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O
target of myo1-like 2 (chicken)
B1 for nudin
olfactory receptor, family 10, subfamily C, member 1
hypothetical protein MGC4276 similar to CG8198

hypothetical protein MGC2776
 chromodomain helicase DNA binding protein 1-like
 Consensus includes gb:BF347113 /FEA=EST /DB_XREF=gi:1294708 /DB_XREF=est:602020879F /CLONE=IMAGE:4156602 /UG=Hs.273138 ESTs
 KIAA1084 protein
 synaptobrevin 2
 stomal antigen 3
 amiloride-sensitive cation channel 2, neuronal
 secretory granule, neuroendocrine protein 1 (782 protein)
 prostate epithelium-specific Ets transcription factor
 endothelial differentiation, sphingolipid G-protein-coupled receptor, 8
 collagen, type V, alpha 2
 KIAA0335 protein
 vsmh-like 1
 KIAA0185 protein
 apolipoprotein H (beta-2-glycoprotein I)
 aradine homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)
 Consensus includes gb:AA293502 /FEA=EST /DB_XREF=gi:1941036 /DB_XREF=est:zt53n06.r1 /CLONE=IMAGE:726107 /UG=Hs.42733 CHMP1.5 protein /FL=gb:AF281
 ATP-binding cassette, sub-family C (CFTR/MRP), member 3
 caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
 four and a half LIM domains 1
 hypothetical protein FLJ11004
 transforming growth factor beta-activated kinase-binding protein 1
 WNT1 inducible signaling pathway protein 1
 EGF-containing fibulin-like extracellular matrix protein 1
 dynein, axonemal, heavy polypeptide 9
 serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antipapain, pigment epithelium derived factor), member 1
 HLA-B associated transcript 8
 Consensus includes gb:AW851086 /FEA=EST /DB_XREF=gi:7946593 /DB_XREF=est:IL3-CT0220-160200-070-802 /UG=Hs.314334 ESTs
 HIV-1 inducer of short transcripts binding protein: lymphoma related factor
 retinoid acid receptor, gamma
 potassium voltage-gated channel, shaker-related subfamily, beta member 1
 Consensus includes gb:A097640 /FEA=EST /DB_XREF=gi:3445898 /DB_XREF=est:qb59a08.x1 /CLONE=IMAGE:1704374 /UG=Hs.66187 Homo sapiens clone 23700 nt
 chromosome 6 open reading frame 32
 high-mobility group (nonhistone chromosomal) protein 1
 tripartite motif-containing 14
 hypothetical protein FLJ13868
 KIAA0322 protein
 peptidylprolyl isomerase (cyclophilin)-like 2
 solute carrier family 21 (organic anion transporter), member 12
 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 1
 low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 8
 peptidylprolyl isomerase (cyclophilin)-like 2
 tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)
 ecdonoidecyl pyrophosphatase/phosphodiesterase 1
 KIAA0582 protein
 Consensus includes gb:A160128 /FEA=EST /DB_XREF=gi:3693508 /DB_XREF=est:qb51n06.x1 /CLONE=IMAGE:1703879 /UG=Hs.295901 KIAA0493 protein
 paired box gene 8
 tenascin XB
 hypothetical protein FLJ11328
 TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor, 135 kD
 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2 (mitochondrial)
 hypothetical protein FLJ20559
 potassium inwardly-rectifying channel, subfamily J, inhibitor 1
 prolins-1-rich protein Bst1 subfamily 1
 DnaJ (Hsp40) homolog, subfamily A, member 4
 Consensus includes gb:AL049991.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G222 (from clone DKFZp564G222) /FEA=mRNA /DB_XREF=gi:4864243 /UG=Hs.2281 AL049991
 neprin A, alpha (PABA peptide hydrolase)
 hypothetical protein DKFZp434P144
 killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 5
 farnesyltransferase, CAAAX box, beta
 zinc finger RNA binding protein
 interferon, alpha 7

Consensus includes gb:AK021633.1 /DEF=Homo sapiens cDNA FLJ11371 fls, clone HEMBA 1003328 /FEA=mRNA /DB_XREF=gi:10432853 /UG=Hs.306608 Homo sapiens AK021633
 Consensus includes gb:AK026449.1 /DEF=Homo sapiens cDNA FLJ22798 fls, clone KAA2544 /FEA=mRNA /DB_XREF=gi:10439319 /UG=Hs.306850 Homo sapiens cdi AK026449
 gb:NM_024118.1 /DEF=Homo sapiens hypothetical protein MGC4692 (MGC4692), mRNA /FEA=mRNA /GEN=MGC4692 /PROD=hypothetical protein MGC4692 /DB_XREF=gi:10439319
 Consensus includes gb:L23852.1 /DEF=Homo sapiens (clone Z146) retinal mRNA, 3' end and repeat region /FEA=mRNA /DB_XREF=gi:393126 /UG=Hs.73838 Homo sapiens
 cyclin-dependent kinase inhibitor 1A (p21, Cip1)
 hypothetical protein FLJ13154
 Falcioni anemia, complementation group E
 Cluster incl. A057637:cy31h06.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-1687483 /clone_end=3 /gb=A057637 /gi=3331503 /ug=Hs.21305 /len=598
 synaptonemal-associated protein, 29kD
 gb:NM_016421.1 /DEF=Homo sapiens relaxininsulin-like factor 1 (LOC51746), mRNA /FEA=CDS /GEN=LOC51746 /PROD=relaxininsulin-like factor 1 /DB_XREF=gi:7708
 KIAA0923 protein
 transcription factor similar to D. melanogaster homeodomain protein lady bird late
 hypothetical protein FLJ11222
 hypothetical protein FLJ14100
 eIF4E transporter
 T cell receptor gamma locus
 activating transcription factor 4 (tax-responsive enhancer element 867)
 heme oxygenase (decycling) 2
 SELENOPHOSPHATE SYNTHETASE : Human selenium donor protein
 serine/threonine kinase 38
 ubiquitin specific protease 4 (proto-oncogene)
 gb:AF095784.1 /DEF=Homo sapiens GABA-B receptor R2 (GABBR2) mRNA, complete cds. /FEA=mRNA /GEN=GABBR2 /PROD=GABA-B receptor R2 /DB_XREF=gi:483
 chorionic somatomammotropin hormone-like 1
 regulator of G-protein signalling 7
 neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)
 hypothetical protein FLJ20894
 v-jun sarcoma virus 17 oncogene homolog (vjun)
 Consensus includes gb:F10172 /FEA=EST /DB_XREF=gi:582615 /DB_XREF=est:hp30h08.x1 /CLONE=IMAGE-271919 /UG=Hs.314230 ESTs, Highly similar to clocd AW238724
 receptor-associated protein of the synapse, 43kD
 KIAA0878 protein
 solute carrier family 17 (sodium phosphate), member 4
 hypothetical protein FLJ21859
 HSPC133 protein
 solute carrier family 16 (monocarboxylic acid transporters), member 5
 CBF1 interacting corepressor
 sarcosine dehydrogenase
 bridging integrator 1
 Consensus includes gb:AW238724 /FEA=EST /DB_XREF=gi:5571113 /DB_XREF=est:hp30h08.x1 /CLONE=IMAGE-271919 /UG=Hs.314230 ESTs, Highly similar to clocd AW238724
 centromere protein F (350/400kD, milstin)
 poliovirus receptor
 hypothetical protein
 colony stimulating factor 1 (macrophage)
 putative UDP-GalNAc:polypeptide N-acetylglucosaminyltransferase T9
 lymphocyte antigen 84 homolog, radioprotective 105kD (mouse)
 oral-facial-digital syndrome 1
 Hermansky-Pudlak syndrome
 protein phosphatase 1, regulatory subunit 10
 transmembrane protease, serine 3
 dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS208, DXS230, DXS238, DXS269, DXS270, DXS272
 hypothetical protein FLJ22593
 integrin, alpha 7
 G6C protein
 interferon (alpha, beta and omega) receptor 1
 cell growth regulatory with ring finger domain
 Fc fragment of IgG, low affinity IIa, receptor for (CD32)
 hypothetical protein, estradiol-induced
 interleukin 15
 single-stranded DNA-binding protein 1
 KIAA0527 protein
 inhibin, beta C
 ATP-binding cassette, sub-family C (CFTR/MRP), member 3
 myobulatin related protein 8
 hypothetical protein FLJ23151

hypothetical protein FLJ21802
 a disintegrin and metalloproteinase domain 20
 leptin (murine obesity homolog)
 LIM domain binding 3
 KIAA0728 protein
 DnaJ (Hsp40) homolog, subfamily B, member 12
 cofactor required for Sp1 transcriptional activation, subunit 2 (150KD)
 granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated serine esterase 3)
 hypothetical protein P1_p373c5
 sin3-associated polypeptide, 30kD
 rhodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4, autosomal dominant)
 hypothetical protein FLJ23323
 Consensus includes gb:AK026465.1 /DEF=Homo sapiens cDNA: FLJ22812 fls, clone KIAU2955, /FEA=mRNA /DB_XREF=gi:10439335 /UG=Hs.306853 Homo sapiens cDNA: AK026465
 hypothetical protein MGC14797
 homolog of mouse C2PA
 tripartite motif-containing 2
 Consensus includes gb:AW451806 /FEA=EST /DB_XREF=gi:5992582 /DB_XREF=est:U1-H-B13-alk-c-10-Q-U1.s1 /CLONE=IMAGE:2737146 /UG=Hs.32458 ESTs
 5-hydroxytryptamine (serotonin) receptor 3A
 secreted protein of unknown function
 solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 6
 lymphocyte antigen 117
 olfactory receptor, family 7, subfamily E, member 47 pseudogene
 gb:BC000233.1 /DEF=Homo sapiens, Similar to guanine nucleotide binding protein, alpha transducing 2, clone MGC:2232, mRNA, complete cds, /FEA=mRNA /PROD=Sin
 ADP-ribosylation factor-like 7
 Consensus includes gb:AK026778.1 /DEF=Homo sapiens cDNA: FLJ23125 fls, clone LNG08217, /FEA=mRNA /DB_XREF=gi:10439709 /UG=Hs.78325 step II splicing factor
 serum amyloid A2
 follistatin
 heparin-binding (tenascin C, cytotactin)
 Consensus includes gb:AV66151.4 /FEA=EST /DB_XREF=gi:3882528 /DB_XREF=est:AV661514 /CLONE=GLCGTC04 /UG=Hs.282693 ESTs
 synaptosomal-associated protein, 23kD
 much 4, tracheobronchial
 KIAA0077 protein
 seroglycan, beta (43kD dystrophin-associated glycoprotein)
 hypothetical protein PRO1866
 UDP-Gal4betaGalNAc beta 1,4'-galactosyltransferase, polypeptide 1
 ADAM-like, decalin 1
 interferon, gamma-inducible protein 16
 Consensus includes gb:BF542055 /FEA=EST /DB_XREF=gi:11629436 /DB_XREF=est:602068320F1 /CLONE=IMAGE:4067357 /UG=Hs.190785 ESTs
 calcium channel, voltage-dependent, L type, alpha 1S subunit
 5'-nucleotidase (putative), cytosolic type B
 vascular Rho-GAP78C-containing
 hypothetical protein FLJ20772
 likely ortholog of mouse O11gp2
 protective protein for beta-galactosidase (galactosialkosis)
 Consensus includes gb:AK021533.1 /DEF=Homo sapiens cDNA: FLJ11471 fls, clone HEMBA1001675, weakly similar to VACUOLAR PROTEIN SORTING-ASSOCIATED P
 CASP8 and FADD-like apoptosis regulator
 nucleoporin 214KD (CAIN)
 ribosomal protein L14
 TONDU
 typtase beta 2
 glyceraldehyde-3-phosphate dehydrogenase, testis-specific
 aconitase 2, mitochondrial
 melanin, ring finger protein, 3
 leucine zipper, putative tumor suppressor 1
 leukocyte-associated Ig-like receptor 1
 guanine nucleotide binding protein (G protein), beta polypeptide 3
 Consensus includes gb:AK025041.1 /DEF=Homo sapiens cDNA: FLJ21388 fls, clone COL03424, /FEA=mRNA /DB_XREF=gi:10437478 /UG=Hs.306751 Homo sapiens cDNA: AK025041
 Consensus includes gb:AT39005 /FEA=EST /DB_XREF=gi:5100986 /DB_XREF=est:w34b03.x1 /CLONE=IMAGE:2392109 /UG=Hs.173448 KIAA0418 gene product
 hypothetical protein FLJ00002
 protein arginine N-methyltransferase 3(hnRNP methyltransferase S. cerevisiae)-like 3
 testis-specific kinase substrate
 Consensus includes gb:AK026889.1 /DEF=Homo sapiens cDNA: FLJ23236 fls, clone COL00725, /FEA=mRNA /DB_XREF=gi:10439654 /UG=Hs.288786 Homo sapiens cDNA: AK026889
 pre-mRNA splicing factor 17
 BG528818

Consensus includes gb:AL049988.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564F212 (from clone DKFZp564F212). /FEA=mRNA /DB_XREF=gi:4884239 /UG=Hs.30631 AL049988
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4
eukaryotic translation elongation factor 1 epsilon 1
eyes absent homolog 4 (Drosophila)
Consensus includes gb:AK025114.1 /DEF=Homo sapiens cDNA: FLJ21461 fls, clone COL04727. /FEA=mRNA /DB_XREF=gi:10437566 /UG=Hs.306768 Homo sapiens cDNA
DKFZP434N014 protein
silver homolog (mouse)
KIAA0633 protein
cell division cycle 27
DKFZP586N0619 protein
KIAA0893 protein
methylthadenosine phosphorylase
leptin receptor
Consensus includes gb:AL137435.1 /DEF=Homo sapiens mRNA: cDNA DKFZp761G0924 (from clone DKFZp761G0924). /FEA=mRNA /DB_XREF=gi:6807994 /UG=Hs.27 AL137435
complement component 5
KIAA0665 gene product
Consensus includes gb:AK024548.1 /DEF=Homo sapiens cDNA: FLJ20895 fls, clone ADKAQ3483. /FEA=mRNA /DB_XREF=gi:10436853 /UG=Hs.285737 Homo sapiens cDNA
transmembrane gamma-carboxyglutamic acid protein 3
latent transforming growth factor beta binding protein 4
Chediak-Higashi syndrome 1
linker for activation of T cells
hypothetical protein PRO1316
centromere protein E (312kD)
fibroblast growth factor 21
sulfotransferase family, cytosolic, 1B, member 1
ribonuclease P (30kD)
engrailed homolog 2
KIAA1053 protein
H4 histone family, member H
likely homolog of mouse immunity-associated nucleotide 4
potassium large conductance calcium-activated channel, subfamily M, alpha member 1
hypothetical protein MGCS627
hydroxyprostaglandin dehydrogenase 15-(NAD)
Consensus includes gb:N35922 /FEA=EST /DB_XREF=gi:1157064 /DB_XREF=est:Y28905.s1 /CLONE=IMAGE:272600 /UG=Hs.269852 ESTs, Weakly similar to ALU1_H N35922
Consensus includes gb:AL046979 /FEA=EST /DB_XREF=gi:5435035 /DB_XREF=est:DKFZp586K0617 /UG=Hs.9973 tennin
neuregulin 1
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3
protease, serine, 1 (trypsin 1)
Consensus includes gb:AU147698 /FEA=EST /DB_XREF=gi:11009219 /DB_XREF=est:AU147698 /CLONE=MAMMA1001408 /UG=Hs.306833 Homo sapiens cDNA FLJ12
Consensus includes gb:BF131965 /FEA=EST /DB_XREF=gi:10971005 /DB_XREF=est:b01820948F1 /CLONE=IMAGE:4052867 /UG=Hs.10684 Homo sapiens clone 2442 BF131965
TIA1 cytotoxic granule-associated RNA binding protein
hypothetical protein FLJ13158
homolog of mouse BMP-2 inducible kinase
hypothetical protein
Consensus includes gb:AA114843 /FEA=EST /DB_XREF=gi:1665911 /DB_XREF=est:20212.s1 /CLONE=IMAGE:491183 /UG=Hs.183454 Homo sapiens mRNA: cDNA DI AA114843
myelin basic protein
lymphoid enhancer-binding factor 1
cellular zinc finger anti-NF-kappaB Cezanne
desmocollin 1
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 2
chorionic gonadotropin hormone 2
taste receptor, type 2, member 3
sialidase 2 (cytosolic sialidase)
SKIP (for skeletal muscle and kidney enriched inositol phosphatase)
splicing factor, arginine/serine-rich 11
B-cell CLL lymphoma 11A (zinc finger protein)
Consensus includes gb:AL049279.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564I083 (from clone DKFZp564I083). /FEA=mRNA /DB_XREF=gi:5000035 /UG=Hs.202886 AL049279
Cdc42 effector protein 3
Rag D protein
protocadherin alpha 8
found in inflammatory zone 3
chromosome 14 open reading frame 2
transcriptional adaptor 2 (ADA2 homolog, yeast)-like

integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
 runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)
 melanoma-associated chondroitin sulfate proteoglycan-like
 protein tyrosine phosphatase, non-receptor type substrate 1
 tumor susceptibility gene 101
 protein tyrosine phosphatase, receptor-type, Z polypeptide 1
 enhancer of zeste homolog 1 (Drosophila)
 KIAA0391 gene product
 tumor necrosis factor receptor superfamily, member 5
 leish-like 3 (Drosophila)
 gp.M33376.1/DEF-Human pseudo-chloride reductase mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:187444 /FL=gb.M33376.1
 WD40 repeat domain 11 protein
 KIAA1105 protein
 transcription factor AP-2, alpha (activating enhancer binding protein 2 alpha)
 cathepsin O
 hyperpolarization activated cyclic nucleotide-gated potassium channel 2
 fibroblast growth factor 18
 Consensus includes gb:AL050065.1 /DEF=Homo sapiens mRNA; cDNA DKFZ566M043 (from clone DKFZ566M043). /FEA=mRNA /DB_XREF=gi:4884295 /UG=Hs.2122 AL050065
 hypothetical protein FLJ20151
 CK2 interacting protein 1; HQ0024c protein
 HBST-like (S. cerevisiae)
 hypothetical protein FLJ13102
 S100 calcium binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11))
 hypothetical protein MG3113
 SH2-B homolog
 activated p21cdc42Hs kinase
 regulator of mitotic spindle assembly 1
 calcium/calmodulin-dependent protein kinase (CaM kinase) II beta
 hypothetical protein FLJ10696
 aquaporin 3
 muscle RAS oncogene homolog
 Consensus includes gb:AL0206718 /FEA=EST /DB_XREF=gi:3765390 /DB_XREF=est:qf61b12.x1 /CLONE=IMAGE:1754495 /UG=Hs.143920 Homo sapiens DNA from chro
 MAD, mothers against decapentaplegic homolog 3 (Drosophila)
 myotubularin related protein 6
 sirin, silent mating type information regulation 2 homolog 3 (S. cerevisiae)
 MAX dimerization protein
 hypothetical protein FLJ10232
 chromosome 9 open reading frame 3
 Huntingtin interacting protein H
 collagen, type XIX, alpha 1
 KIAA1243 protein
 dual specificity phosphatase 8
 hepatocellular carcinoma-associated antigen 112
 Consensus includes gb:BF593594 /FEA=EST /DB_XREF=gi:11685918 /DB_XREF=est:q12a08.x1 /CLONE=IMAGE:3697838 /UG=Hs.274440 Homo sapiens mRNA; cDN
 chaperonin containing TCP1, subunit 8A (zeta 1)
 Consensus includes gb:AL031651 /DEF=Human DNA sequence from clone RP5-1054A22 on chromosome 20q11.22-12 Contains two isoforms of the gene for TGM2 (trans
 nuclear receptor subfamily 2, group C, member 1
 integrin, alpha 6
 calixin
 Cluster Inc. A1970898:wr1cd3.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2488324 /clone_and=3 /gb=A1970898 /gi=5767774 /ug=Hs.234698 /len=382
 thoredoxin-like, 32KD
 olfactory receptor, family 7, subfamily E, member 12 pseudogene
 single-minded homolog 2 (Drosophila)
 adenylate cyclase 6
 cyclic AMP phosphoprotein, 19 kD
 Consensus includes gb:A1879213 /FEA=EST /DB_XREF=gi:4869395 /DB_XREF=est:hw72p04.x1 /CLONE=IMAGE:2256630 /UG=Hs.81920 ESTs, Weakly similar to cDNA I
 BTB and CNC homolog 1, basic leucine zipper transcription factor 1
 hypothetical protein
 hypothetical protein FLJ10055
 carnitine palmitoyltransferase 1, liver
 transcription factor Dp-2 (E2F dimerization partner 2)
 paired basic amino acid cleaving system 4
 hypothetical protein FLJ10738

baculoviral IAP repeat-containing 4
 immunoglobulin lambda variable (V)OR22-1
 hypothetical protein FLJ22233
 pre-mRNA processing factor 31 homolog (yeast)
 gb:M32577.1 /DEF=Human MHC HLA-DQ beta mRNA, complete cds. /FEA=mRNA /GEN=HLA-DQ81 /DB_XREF=gi:188194 /FL=gb:M32577.1
 DKFZP564B147 protein
 NIF3 NGG1 interacting factor 3-like 1 (S. pombe)
 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced)
 hypothetical protein DKFZp434L0718
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 5
 cAMP responsive element modulator
 stannocalcin 1
 diacylglycerol kinase, alpha (80kD)
 hypothetical protein FLJ10349
 keratin 6A
 Interleukin 1 receptor accessory protein
 (clone PWHL22-24) myosin light chain 2
 hypothetical protein FLJ10388
 endometrial bleeding associated factor (left-right determination), factor A; transforming growth factor beta superfamily)
 transforming, acidic coiled-coil containing protein, 3
 tropomyosin 2 (beta)
 Consensus includes gb:AL157398 /DEF=Human DNA sequence from clone RP11-56H7 on chromosome 10. Contains ESTs, STSs and GSSs. Contains the gene for the re
 HIV-1 Rev binding protein
 hypothetical protein FLJ10856
 von Willebrand factor
 Consensus includes gb:AU145354 /FEA=EST /DB_XREF=gi:11006875 /DB_XREF=est:AU145354 /CLONE=HEMBA1004608 /UG=Hs.287411 Homo sapiens cDNA FLJ111
 KIAA0597 protein
 mannan-binding lectin serine protease 1 (CA/C2 activating component of Rf-reactive factor)
 CCAAT/enhancer binding protein (C/EBP), alpha
 zinc finger protein 148 (pH2-52)
 ELK1, member of ETS oncogene family
 KIAA0248 protein
 Ras-related GTP-binding protein
 son of sevenless (Drosophila) homolog 2
 Cluster incl. AA524093.ng3304.s1 Homo sapiens cDNA, 3 end /clone=IMAGE:936607 /clone_end=3 /gb=AA524093 /gi=2265021 /ug=Hs.23158 /len=718
 erythropoietin receptor
 H1 histone family, member 0
 glycerol kinase
 solute carrier family 12 (sodium/potassium/chloride transporters), member 2
 activating transcription factor 2
 cystinosis, nephropathic
 transcription factor-like 1
 ribonuclease, RNase A family, 4
 hypothetical protein MGC2849
 tyrosinase beta 1
 naturalistic peptide precursor B
 hypothetical protein
 RNA binding motif protein 9
 ems1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate)
 insulin-like growth factor 2 receptor
 Consensus includes gb:AK024525.1 /DEF=Homo sapiens cDNA: FLJ20872 fls, clone AOKA02804. /FEA=mRNA /DB_XREF=gi:10436827 /UG=Hs.306683 Homo sapiens
 neuromedin B
 protein phosphatase 2A, regulatory subunit B' (PR 53)
 fls485
 karyopherin beta 2b, transportin
 hypothetical protein FLJ10579
 ubiquitin activating enzyme E1-like protein
 nescent helix loop helix 2
 gap junction protein, alpha 3, -46kD (connexin 46)
 hypothetical protein FLJ13188
 eukaryotic translation initiation factor 4E-like 3
 Consensus includes gb:AL050053.1 /DEF=Homo sapiens mRNA: cDNA DKFZp566E103 (from clone DKFZp566E103) /FEA=mRNA /DB_XREF=gi:4864289 /UG=Hs.2273:AL050053
 KIAA0793 gene product
 NM_001167 1.05
 AL008721 1.05
 NM_024859 1.05
 NM_015628 1.05
 M32577 1.05
 NM_015582 1.05
 NM_021824 1.05
 BC005254 1.05
 AL136784 1.05
 NM_002639 1.05
 S68134 1.05
 AL300520 1.05
 AF064771 1.05
 NM_018086 1.05
 AL569511 1.05
 NM_002182 1.05
 NM_013292 1.05
 NM_018082 1.05
 NM_003240 1.05
 NM_006342 1.05
 NM_003269 1.05
 NM_0157398 1.05
 A742626 1.05
 NM_018247 1.05
 NM_000552 1.05
 NM_145354 1.05
 A656706 1.05
 NM_001879 1.05
 NM_004364 1.05
 NM_021964 1.05
 AF000872 1.05
 D87433 1.05
 NM_006570 1.05
 A1276593 1.05
 AA524093 1.05
 M34988 1.05
 Z97630 1.05
 A830490 1.05
 NM_001048 1.05
 NM_001880 1.05
 AJ222967 1.05
 NM_005997 1.05
 A1781728 1.05
 NM_024080 1.04
 AF206667 1.04
 NM_002521 1.04
 AF151074 1.04
 N95026 1.04
 NM_005231 1.04
 BG031974 1.04
 NM_021077 1.04
 X86428 1.04
 NM_015931 1.04
 BG256639 1.04
 NM_018145 1.04
 NM_006395 1.04
 AA166895 1.04
 NM_021954 1.04
 NM_022063 1.04
 BF516289 1.04
 NM_014508 1.04

ataxin 2 related protein	NM_007245	1.04
KIAA1882 protein	BC003850	1.04
SPL protein	NM_018269	1.04
cydin D2	AW026491	1.04
hypothetical protein FLJ22187	NM_024533	1.04
hypothetical protein DKFZp434A1022	NM_021185	1.04
histone acetyltransferase 1	NM_003642	1.04
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	NM_005228	1.04
Consensus includes gb:R61519 /FEA=EST /DB_XREF=gi:302214 /DB_XREF=est:YH1804.s1 /CLONE=IMAGE:37896 /UG=Hs.185807 Homo sapiens done 24758 mRNA	R61519	1.04
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	U08015	1.04
calcium/calmodulin-dependent protein kinase kinase 2, beta	AA181179	1.04
erythropoietin receptor	M60459	1.04
modulin	NM_002418	1.04
1003 protein	NM_020232	1.04
disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)	AF188298	1.04
myosin, light polypeptide 3, alkali, ventricular, skeletal, slow	NM_000258	1.04
cytochrome P450 polypeptide 43	AF280110	1.04
ets variant gene 5 (ets-related molecule)	NM_004454	1.04
cadherin 7, type 2	NM_004381	1.04
glutathione S-transferase A3	NM_000847	1.04
Consensus includes gb:AL080182.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O151 (from clone DKFZp434O151) /FEA=mRNA /DB_XREF=gi:5262658 /UG=Hs.2251	AL080182	1.04
a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	U86755	1.04
plasminogen activator, urokinase receptor	AY029180	1.04
chloride channel 3	AJ760629	1.04
hypothetical protein FLJ20707	NM_017936	1.04
UV-B repressed sequence, HUR 7	BE148534	1.04
glutathione S-transferase subunit 13 homolog	NM_015917	1.04
clone FLB3816	NM_016415	1.04
KIAA1048 protein	AW451954	1.04
serologically defined colon cancer antigen 16	NM_006649	1.04
zinc finger protein, subfamily 1A, 2 (Helios)	NM_018280	1.04
monocamine oxidase A	AA923354	1.04
hypothetical protein MGC5356	NM_024059	1.04
PRO0214 protein	NM_014120	1.04
Consensus includes gb:AK024873.1 /DEF=Homo sapiens cDNA; FLJ21220 fts, clone COL00546 /FEA=mRNA /DB_XREF=gi:10437285 /UG=Hs.306713 Homo sapiens cDNA	AK024873	1.04
Fancconi anemia, complementation group C	NM_000136	1.04
transcription factor 8 (represses interleukin 2 expression)	AI373166	1.04
hypothetical protein FLJ22175	AA551370	1.04
nuclear mitotic apparatus protein 1	AI337584	1.04
hypothetical protein MGC3123	NM_024107	1.04
KIAA0862 protein	AK022530	1.04
paralemmin	NM_002579	1.04
hypothetical protein FLJ12838	NM_024641	1.04
dectand homolog (Drosophila)	NM_004392	1.04
hypothetical protein FLJ23229	NM_025058	1.04
insulin-like growth factor 1 (somatomedin C)	M25644	1.04
adenosine A2b receptor pseudogene	AL022398	1.04
N-myc downstream-regulated gene 3	BC001805	1.04
Consensus includes gb:BG150564 /FEA=EST /DB_XREF=gi:12662594 /DB_XREF=est:K02G04.x1 /CLONE=IMAGE:3443143 /UG=Hs.184326 CDC10 (cell division cycle	BC150564	1.04
inosine triphosphatase (nucleoside triphosphate pyrophosphatase)	AF219116	1.04
calcium channel, voltage-dependent, alpha 2/delta 3 subunit	NM_000277	1.04
KIAA0578 protein	NM_018398	1.04
hypothetical protein	AU144775	1.04
adaptor-related protein complex 1, sigma 1 subunit	NM_019073	1.04
small inducible cytokine subfamily A (Cys-Cys), member 11 (eotaxin)	NM_001283	1.04
formyltetrahydrofolate dehydrogenase	D49372	1.04
EphB2	NM_012190	1.04
progesterone receptor	D31681	1.04
paired-like homeodomain transcription factor 3	NM_000926	1.04
phosphatidylesterase-3-kinase, regulatory subunit, polypeptide 3 (p55, gamma)	NM_005029	1.04
hypothetical protein FLJ11818	AF028785	1.04
zinc finger protein 238	NM_022452	1.04
	NM_006352	1.04

KIAA0427 gene product	U55962	1.04
leukocyte-associated Ig-like receptor 1	NM_021708	1.04
KIAA0261 protein	AK027005	1.04
hypothetical protein	NM_015689	1.04
TGF-beta-induced factor 2 (TALE family homeobox)	NM_021809	1.04
bladder cancer associated protein	NM_006898	1.04
hypothetical protein FLJ11585	AI743854	1.04
glutamate receptor, ionotropic, N-methyl D-aspartate 2D	NM_000836	1.04
DKFZP586P2219 protein	W80642	1.04
KIAA0717 protein	AY009093	1.04
endocrine regulator	AW613549	1.04
nuclear autoantigen (SSKd) similar to rat synaptonemal complex protein	NM_006455	1.04
aldohyde dehydrogenase 1 family, member A1	NM_000889	1.04
elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1	NM_016031	1.04
homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2	AL136939	1.04
kinesin family member 4A	NM_012310	1.04
tol-like receptor 4	NM_003266	1.04
hypothetical protein FLJ20184	NM_017700	1.04
exosome component Rps41	NM_018037	1.04
hypothetical protein FLJ20279	NM_017740	1.04
Consensus includes gb:AB007970.1 /DEF=Homo sapiens mRNA, chromosome 1 specific transcript KIAA0501. /FEA=mRNA /DB_XREF=gi:3413945 /UG=Hs.223020 Homt	AB007970	1.04
chorionic gonadotropin hormone 1 (placental lactogen)	BC005921	1.04
signal transducer and activator of transcription 5B	NM_012448	1.04
hypothetical protein FLJ21007	NM_030794	1.04
Consensus includes gb:A1895595 /FEA=EST /DB_XREF=gi:4983495 /DB_XREF=est:we49g06.x1 /CLONE=IMAGE:2344474 /UG=Hs.293219 ESTs	A1895595	1.04
coagulation factor III (thromboplastin, tissue factor)	NM_001993	1.04
proliferation factor 13	NM_002818	1.04
Consensus includes gb:AK024177.1 /DEF=Homo sapiens cDNA FLJ14115 fls. clone MAMMA1001760. /FEA=mRNA /DB_XREF=gi:10436493 /UG=Hs.266263 Homo sapie	AK024177	1.04
hypothetical protein Z4636	AU150186	1.04
promyelocytic leukemia	M79462	1.04
Consensus includes gb:L34409.1 /DEF=Homo Sapiens (clone B383E13) chromosome 4p16.3 DNA fragment. /FEA=mRNA /DB_XREF=gi:800517 /UG=Hs.94799 Homo St	L34409	1.04
proline arginine-rich and leucine-rich repeat protein	NM_002725	1.04
MAD2 mitotic arrest deficient-like 1 (yeast)	NM_002358	1.04
putative DNA binding protein	AF072814	1.04
tyrosine aminotransferase	AV647713	1.04
KIAA0342 gene product	AQ035414	1.04
wingless-type MMTV integration site family, member 16	NM_016087	1.04
MAD, mothers against decapentaplegic homolog 3 (Drosophila)	NM_005902	1.04
growth suppressor 1	NM_022356	1.04
KIAA1087 protein	AB028980	1.04
cytochrome P450, subfamily IIIA (naphthalene oxidase), polypeptide 5	NM_000777	1.04
promyelocytic leukemia	NM_002675	1.04
G protein-coupled receptor kinase 6	AF040752	1.04
N-acylphosphatase amidohydrolase (acid ceramidase)	A1934569	1.04
Consensus includes gb:AL133519 /DEF=Human DNA sequence from clone RP11-12201 on chromosome 20 Contains a novel gene encoding a protein orthologous to the	AL133519	1.04
PKS1 postmeiotic segregation increased 1 (S. cerevisiae)	BG434893	1.04
Consensus includes gb:AW081113 /FEA=EST /DB_XREF=gi:6036265 /DB_XREF=est:xc29c08.x1 /CLONE=IMAGE:2585678 /UG=Hs.18368 DKFZP56480789 protein	AW081113	1.04
CGI-86 protein	AF126782	1.04
mitogen-activated protein kinase kinase kinase 4	NM_004834	1.04
small nuclear ribonucleoprotein polypeptide E	NM_003094	1.04
Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF=AF005082	AF005082	1.04
hypothetical protein p5326	AF073483	1.04
cyclin-dependent kinase 8	R59697	1.04
KIAA1074 protein	NM_014915	1.04
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	Y09216	1.04
E74-like factor 2 (eta domain transcription factor)	NM_006874	1.04
tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain	AF012538	1.04
block of proliferation 1	AK024840	1.04
Consensus includes gb:AL137325.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M0835 (from clone DKFZp434M0835) /FEA=mRNA /DB_XREF=gi:6807816 /UG=Hs.27 AL137325	AL137325	1.04
SON DNA binding protein	NM_003103	1.04
basic helix-loop-helix domain containing, class B, 2	BG326045	1.04
uncharacterized hypothalamus protein HT012	NM_018473	1.04
nidogen 2	NM_007361	1.04

323

cystinosis, nephropathic	NM_004937	1.04
zuculin related factor 1	AI338837	1.04
secretory pathway component Sec31B-1	AF274863	1.04
Neuro-d4 (rat) homolog	NM_004647	1.04
myosin, light polypeptide 1, alkali; skeletal, fast	M20643	1.04
fatty acid amide hydrolase	NM_001441	1.04
protein similar to E.coli yhdg and R. capsulatus nifR3	NM_007016	1.04
lipase protein	AF225418	1.04
glutamate receptor, ionotropic, AMPA 4	NM_000829	1.04
ligase III, DNA, ATP-dependent	NM_013975	1.04
hypothetical protein MGC2745	NM_024117	1.04
collagen, type VI, alpha 2	AL531750	1.04
regulator of G-protein signalling 1	SS9049	1.04
Consensus includes gb:AW663888 /FEA=EST /DB_XREF=gi:7456424 /DB_XREF=est:ht73602.x1 /CLONE=IMAGE:2977699 /UG=Hs.43834 Homo sapiens immunoglobulin	AFV663885	1.04
breifeldin A-inhibited guanine nucleotide-exchange protein 1	AF084320	1.04
calcium channel, voltage-dependent, gamma subunit 2	AL022313	1.04
major histocompatibility complex, class II, DR beta 4	BC005312	1.04
fused leuc homolog (mouse)	NM_022476	1.04
nuclear receptor coactivator 3	AF010227	1.04
calcineurin binding protein 1	NM_012295	1.04
prostate epithelium-specific Ets transcription factor	NM_012391	1.04
acrosomal vesicle protein 1	NM_020113	1.04
potassium voltage-gated channel, subfamily H (eag-related), member 2	AB044806	1.04
dual specificity phosphatase 7	X93921	1.04
peroxisome proliferative activated receptor, delta	L07592	1.04
tumor necrosis factor receptor superfamily, member 6b, decoy	NM_003823	1.04
absent in melanoma 1	U83115	1.04
vesicle-associated membrane protein 5 (myobrevin)	AB14466	1.04
zinc finger protein 202	N81520	1.04
serine/threonine kinase 228 (spemiotogenesis associated)	AB52441	1.04
tousted-like kinase 1	NM_012290	1.04
Consensus includes gb:AW517688 /FEA=EST /DB_XREF=gi:7155768 /DB_XREF=est:xs86405.x1 /CLONE=IMAGE:2776521 /UG=Hs.305923 ATPase, Ca++ transporting,	AW517686	1.04
nodulin 2	NM_004475	1.04
makorin, ring finger protein, 1	NM_013446	1.04
myobulbin related protein 2	NM_016156	1.04
potassium voltage-gated channel, Isk-related family, member 1	NM_000219	1.04
hypothetical protein FLJ20342	NM_017774	1.04
hypothetical protein FLJ22729	NM_024683	1.04
polymerase (DNA directed), gamma 2, accessory subunit	NM_007215	1.04
KIAA0980 protein	NM_025176	1.04
chromosome 1 open reading frame 28	NM_024529	1.04
protein tyrosine phosphatase, non-receptor type 21	BF116199	1.04
discs, large (Drosophila) homolog 1	AW139131	1.04
non-metastatic cells 3, protein expressed in	AL031718	1.04
latent transforming growth factor beta binding protein 1	NM_006627	1.04
ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle	AU140931	1.04
hemochromatosis	AF144244	1.04
dimethylarginine dimethylaminohydrolase 2	NM_013974	1.04
Consensus includes gb:AI692428 /FEA=EST /DB_XREF=gi:4969768 /DB_XREF=est:wd69c12.x1 /CLONE=IMAGE:2336854 /UG=Hs.184019 Homo sapiens clone 23551	AI692428	1.04
bile acid Coenzyme A: amino acid N-acyltransferase (glycine N-choloyltransferase)	NM_001701	1.04
DIPB protein	BF431488	1.04
serine/glucocorticoid regulated kinase	NM_005827	1.04
BET1 homolog (S. cerevisiae)	BC000899	1.04
TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31 kD	AF077053	1.04
immunoglobulin kappa constant	X72475	1.04
RAO31-interacting protein	BE966146	1.04
ecodonucleoside triphosphate diphosphohydrolase 1	U87967	1.04
retinal outer segment membrane protein 1	NM_000327	1.04
integrin receptor matrix proteoglycan 200	NM_016247	1.04
hypothetical protein FLJ21919	NM_023015	1.04
polyglutamine binding protein 1	AB041834	1.04
hypothetical protein FLJ10199	NM_018022	1.04
myeloid leukemia factor 2	NM_005439	1.04

gamma-tubulin complex protein 2	BF002130	1.04
glycine receptor, beta	AF094754	1.04
excision repair cross-complementing rodent repair deficiency, complementation group 4	U64315	1.04
phorbol-like protein MDS019	BF508948	1.04
melanoma antigen, family A, 12	U82671	1.04
hypothetical protein FLJ10483	NM_018108	1.04
KIAA0785 protein	NM_025010	1.04
platin B2	BC004542	1.04
KIAA0785 protein	AB002293	1.04
Consensus includes gb:AA721025 /FEA=EST /DB_XREF=gi:2737160 /DB_XREF=est:nd8905.s1 /CLONE=IMAGE:1269464 /UG=Hs.293253 ESTs	AA721025	1.04
hypothetical protein MY014	BF938727	1.04
tumor necrosis factor, alpha-induced protein 6	AW188198	1.04
KIAA0633 gene product	NM_014645	1.04
PR domain containing 2, with ZNF domain	U23736	1.04
gb:M26463.1 /DEF=Homo sapiens immunoglobulin mu chain antibody MO30 (IgM) mRNA, complete cds. /FEA=CDS /GEN=IgM /PROD=immunoglobulin mu chain antibody	M28463	1.04
prostate epithelium-specific Ets transcription factor	A1307815	1.04
synaptophysin 1	BC000731	1.04
SUMO-1-specific protease	NM_015571	1.04
FYN binding protein (FYB-120/130)	A633888	1.04
hypothetical protein FLJ10901	NM_018265	1.04
cadherin 3, type 1, P-cadherin (placental)	NM_001793	1.04
glutaredoxin (thioltransferase)	NM_002064	1.04
lactoferrin rich protein	BF665706	1.04
zinc finger protein 219	NM_016423	1.04
calcium channel, voltage-dependent, alpha 1G subunit	AF126966	1.04
heparan sulfate proteoglycan 2 (perlecan)	A981033	1.04
Consensus includes gb:AK022309.1 /DEF=Homo sapiens cDNA FLJ12247 fls, clone MAMMA1001397. /FEA=mRNA /DB_XREF=gi:10433678 /UG=Hs.287506 Homo sapiens	AK022309	1.04
KIAA1888 protein	AW444969	1.04
hypothetical protein FLJ12618	NM_024884	1.03
microtubule-associated protein, RPIEB family, member 3	A835178	1.03
nuclear receptor subfamily 2, group C, member 2	NM_003298	1.03
CD209 antigen-like	NM_014257	1.03
Consensus includes gb:AU146794 /FEA=EST /DB_XREF=gi:11008315 /DB_XREF=est:AU146794 /CLONE=HEMBB1001536 /UG=Hs.287475 Homo sapiens cDNA FLJ121	AU146794	1.03
prolactin-induced protein	NM_002652	1.03
procollagen C-endopeptidase enhancer 2	NM_013363	1.03
hypothetical protein FLJ22569	NM_023925	1.03
soluble carrier family 24 (sodium/potassium/calcium exchanger), member 2	NM_020344	1.03
uncharacterized hypothetical protein HCDASE	NM_018479	1.03
CGI-96 protein	AL589666	1.03
H2A histone family, member D	BF061074	1.03
homoeo box A9	U41813	1.03
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	NM_002810	1.03
Consensus includes gb:AL121994 /DEF=Human DNA sequence from clone RP4-781L3 on chromosome 1p34.3-36.11 Contains a pseudogene similar to IFITM3 (interferon	AL121994	1.03
Consensus includes gb:BF222018 /FEA=EST /DB_XREF=gi:1129195 /DB_XREF=est:7p41a04.x1 /CLONE=IMAGE:3648271 /UG=Hs.289603 ESTs. Moderately similar to BF222018	BF222018	1.03
microseminoprotein, beta-	NM_002443	1.03
KIAA1654 protein	AB051441	1.03
bone morphogenetic protein 2	NM_001200	1.03
collagen, type V, alpha 1	M76729	1.03
cytidine monophosphate-N-acetylneuraminic acid hydroxylase (CMP-N-acetylneuraminic acid monooxygenase)	AF074480	1.03
adipose differentiation-related protein	BC005127	1.03
Consensus includes gb:L43577 /FEA=EST /DB_XREF=gi:899066 /DB_XREF=est:L43577 /CLONE=EST500 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43577	1.03
B-cell linker	NM_013314	1.03
KIAA0284 protein	AB006622	1.03
uncharacterized hypothetical protein HT010	NM_018471	1.03
solute carrier family 6 (neurotransmitter transporter, serotonin), member 4	NM_001045	1.03
eosinophil peroxidase	X14346	1.03
Consensus includes gb:A871747 /FEA=EST /DB_XREF=gi:4851478 /DB_XREF=est:wa05e06.x1 /CLONE=IMAGE:2297218 /UG=Hs.81360 Homo sapiens cDNA: FLJ2195; A871747	A871747	1.03
chromosome 21 open reading frame 80	BC000626	1.03
tubby like protein 3	A1591305	1.03
hypothetical protein FLJ20972	NM_025030	1.03
glucosyltransferase 6 (alpha (1,3) glucosyltransferase)	M98625	1.03
Conserved gene telomeric to alpha globin cluster	BE791629	1.03
CGI-01 protein	AL049669	1.03

protein tyrosine phosphatase, receptor type, J	O37781	1.03
peptide YY	D13902	1.03
catenin (cadherin-associated protein), delta 1	NM_001331	1.03
mitochondrial intermediate peptidase	U80034	1.03
TAF4B RNA polymerase II, TATA box binding protein (TBP)-associated factor, 105 MD	U09321	1.03
laminin, alpha 4	U77706	1.03
histatin 3	NM_000200	1.03
receptor tyrosine kinase-like orphan receptor 1	NM_005012	1.03
fibulin 5	NM_006329	1.03
hypothetical gene MGC9753	BF033007	1.03
mannosidase, beta A, lysosomal	NM_005908	1.03
KIAA0977 protein	BF002844	1.03
Consensus includes gb:AW161628 /FEA=EST /DB_XREF=gi:6330659 /DB_XREF=est:au68b11.x1 /CLONE=IMAGE:2781405 /UG=Hs.21739 Homo sapiens mRNA; cDNA	AW161628	1.03
KIAA1354 protein	AW136584	1.03
bone morphogenetic protein 7 (osteogenic protein 1)	AL157414	1.03
chromosome 1 open reading frame 24	NM_022083	1.03
chemokine (C-C motif) receptor 2	NM_000647	1.03
agouti related protein homolog (mouse)	NM_001138	1.03
retinol binding protein 3, interstitial	J03912	1.03
nucleolar protein ANKT	NM_016359	1.03
Kruppel-like factor 2 (lung)	NM_016270	1.03
KIAA1115 protein	BC002789	1.03
caveolin 1, caveolae protein, 22kD	AU147399	1.03
phospholipase D1, phosphatidylcholine-specific	AJ276230	1.03
ATPase, Class V, type 10D	AU78147	1.03
adrenergic, beta-2-, receptor, surface	NM_000024	1.03
hypothetical protein MGC2487	NM_023932	1.03
Consensus includes gb:AU155105 /FEA=EST /DB_XREF=gi:11016626 /DB_XREF=est:AU155105 /CLONE=OVARC:1001000 /UG=Hs.301348 Homo sapiens cDNA FLJ13	NM_023932	1.03
phospholipase A2, group IVB (cytosolic)	NT1116	1.03
wingless-type MMTV integration site family, member 2B	AB045117	1.03
myo48 protein	AF063608	1.03
neuronal cell adhesion molecule	U55258	1.03
ubiquitin-conjugating enzyme E2D 2 (UBC4/S homolog, yeast)	BF196642	1.03
CLIP-associated protein 2	AL137636	1.03
RAS p21 protein activator (GTPase activating protein) 3 (ins1(1.3.4.5)/p4-binding protein)	NM_007368	1.03
Consensus includes gb:AK000144.1 /DEF=Homo sapiens cDNA FLJ20137 fls, clone COL07137. /FEA=mRNA /DB_XREF=gi:7020040 /UG=Hs.274449 Homo sapiens cDN	NM_007368	1.03
solute carrier family 25 (mitochondrial oxalocarboxylate carrier), member 21	NM_030631	1.03
NGF1-A binding protein 1 (EGF1 binding protein 1)	AF045452	1.03
ribophorn II	AL514285	1.03
thyrotropin-releasing hormone receptor	O16845	1.03
PDZ-LIM protein mystique	NM_021630	1.03
centrosomal P4, 1-associated protein	NM_018451	1.03
UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylgalactosaminyltransferase 6 (GalNAc-T6)	NM_007210	1.03
major vault protein	NM_017458	1.03
Sp1 transcription factor	AU121035	1.03
diaphorase (NADH) (cytochrome b-5 reductase)	NM_000398	1.03
signal recognition particle receptor (dockin protein)	BC414541	1.03
nuclear factor (erythroid-derived 2)-like 3	NM_004289	1.03
Consensus includes gb:AW071997 /FEA=EST /DB_XREF=gi:6026995 /DB_XREF=est:ws5709.x1 /CLONE=IMAGE:2501321 /UG=Hs.328248 Homo sapiens mRNA; cDNA	NM_004289	1.03
Consensus includes gb:AF218683 /DEF=Homo sapiens Interleukin-1 receptor antagonist homolog 1 (IL1HY1) gene, complete cds /FEA=mRNA, 1 /DB_XREF=gi:8572054 /	AF218683	1.03
Cluster ind. H59033.yr40e10.r1 Homo sapiens cDNA, 5' end /clone=IMAGE:207786 /clone_end=5' /gb=H59033 /gi=1011865 /ug=Hs.6700 /len=604	H59033	1.03
elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1	H93026	1.03
deafness locus associated putative guanine nucleotide exchange factor	NM_012139	1.03
cell adhesion molecule with homology to L1CAM (close homologue of L1)	NM_006614	1.03
highly expressed in cancer, rich in leucine heptad repeats	NM_006101	1.03
hypothetical protein FLJ11730	NM_022756	1.03
hypothetical protein MGC4342	NM_024329	1.03
hypothetical protein FLJ13189	NM_024682	1.03
KIAA0564 protein	AU744626	1.03
peroxisomal long-chain acyl-coA thioesterase	NM_006821	1.03
adaptor-related protein complex 3, mu 2 subunit	NM_006803	1.03
hypothetical protein MGC11242	NM_024320	1.03
Fc fragment of IgG, low affinity Iib, receptor for (CD32)	U09040	1.03

Consensus includes gb:AW17304.5 /FEA=EST /DB_XREF=gi:6438993 /DB_XREF=est:82603.x1 /CLONE=IMAGE:2663740 /UG=Hs.322856 Homo sapiens cDNA FLJ101 AW17304.5
 mannosidase, alpha, class 2A, member 1
 1.03 NM_002372
 1.03 NM_001884
 cartilage linking protein 1
 1.03 D13705
 Consensus includes gb:BG387770 /FEA=EST /DB_XREF=gi:13281216 /DB_XREF=est:602412648F1 /CLONE=IMAGE:4521442 /UG=Hs.147025 ESTs, Weakly similar to (BG387770
 KIAA0864 protein
 1.03 N31673
 coagulation factor VIII, procoagulant component (hemophilia A)
 1.03 NM_000132
 1.03 NM_002063
 glycoprotein P450, subfamily IVA, polypeptide 11
 1.03 NM_005665
 ecotropic viral integration site 5
 1.03 AW166067
 ribosome binding protein 1 homolog 180kD (dog)
 1.03 NM_004166
 small inducible cytokine subfamily A (Cys-Cys), member 14
 1.03 NM_007351
 multimerin
 1.03 NM_004591
 small inducible cytokine subfamily A (Cys-Cys), member 20
 1.03 AL050220
 kallikrein 13
 1.03 NM_005887
 deleted in lymphocytic leukemia, 1
 1.03 NM_024741
 hypothetical protein FLJ12827
 1.03 U06935
 thyrotrophic embryonic factor
 1.03 A1962943
 hypothetical protein FLJ22028
 1.03 AV713053
 hypothetical protein FLJ14547
 Consensus includes gb:A1133721 /FEA=EST /DB_XREF=gi:6361037 /DB_XREF=est:HA2346 /UG=Hs.53112 ESTs, Moderately similar to ALU8_HUMAN ALU SUBFAMILY A1133721
 1.03 NM_004258
 immunoglobulin superfamily, member 2
 1.03 M26095
 heterologous nuclear ribonucleoprotein A0
 1.03 BE966599
 hypothetical protein MGC10334
 1.03 NM_030575
 KIAA1402 protein
 Consensus includes gb:AA554430 /FEA=EST /DB_XREF=gi:2324969 /DB_XREF=est:n11404.s1 /CLONE=IMAGE:1030303 /UG=Hs.269421 Homo sapiens cDNA FLJ1434 AA554430
 1.03 AB037823
 hypothetical protein FLJ14260
 1.03 NM_025027
 hypothetical protein PRO2405
 1.03 NM_018627
 acylphosphatase 1, erythrocyte (common) type
 1.03 NM_001107
 hypothetical gene FLJ00060
 1.03 AK024467
 gb:NM_002848.2 /DEF=Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO), transcript variant 2, mRNA, FEA=mrna /GEN=PTPRO /PROD=receptor, NM_002848
 KIAA0808 gene product
 1.03 NM_014729
 hypothetical protein FLJ10837
 1.03 AF274950
 uncoupling protein 3 (mitochondrial, proton carrier)
 1.03 NM_022803
 enst1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate)
 1.03 BG475299
 S100 calcium binding protein P
 1.03 NM_005980
 actin-like TA
 1.03 NM_006687
 signal recognition particle 72kD
 1.03 A453872
 hypothetical protein FLJ12229
 1.03 NM_024876
 carnitine acyltransferase
 1.03 BC000723
 hypothetical protein FLJ23182
 1.03 NM_022366
 chloride channel 3
 1.03 AF029346
 hemoglobin, beta
 1.03 M25078
 KIAA1402 protein
 1.03 AA534198
 p65 protein
 1.03 NM_017532
 ZAP3 protein
 1.03 A1703162
 golgi transport complex 1 (80 kDa subunit)
 1.03 AU152134
 hypothetical protein MGC19556
 1.03 AV746402
 Consensus includes gb:AU158442 /FEA=EST /DB_XREF=gi:11019963 /DB_XREF=est:AU158442 /CLONE=PLACE2000394 /UG=Hs.273099 Homo sapiens cDNA FLJ137 AU158442
 1.03 NM_003885
 cyclin-dependent kinase 5, regulatory subunit 1 (p35)
 1.03 NM_001215
 carbonic anhydrase VI
 1.03 AKG25059
 KIAA1332 protein
 1.03 NM_015218
 KIAA0433 protein
 1.03 NM_004530
 matrix metalloproteinase 2 (gelatinase A, 72kD type IV collagenase)
 1.03 BF689173
 DKFZPS86M1523 protein
 1.03 BC000229
 hypothetical protein MGC2488
 1.03 NM_003357
 uteroglobin
 1.03 NM_016603
 chromosome 5 open reading frame 5
 1.03 R38389
 olfactomedin 1
 1.03 NM_001155
 paired immunoglobulin-like receptor alpha
 Consensus includes gb:AK001155.1 /DEF=Homo sapiens cDNA FLJ10293 f1a, clone NT2RM1000280, highly similar to VACUOLAR ATP SYNTHASE SUBUNIT O (EC 3.6.1.4)
 1.03 AK001155
 amyloid beta (A4) precursor protein-binding, family A, member 1 (X11)
 1.03 NM_001163
 ectodermal-neural cortex (with BTB-like domain)
 1.03 AF010314

329

ATP-dependent RNA helicase	AF077033	1.02
synaptonemal complex protein 2	NM_014258	1.02
LDL receptor adaptor protein	AL545035	1.02
KIAA0586 gene product	NM_014749	1.02
KIAA0700 protein	AW450345	1.02
Consensus includes gb:BG105365 /FEA=EST /DB_XREF=gi:12599141 /DB_XREF=est:602312473F1 /CLONE=IMAGE:4422043 /UG=Hs.23348 S-phase kinase-associated	BG105365	1.02
Consensus includes gb:AK023365.1 /DEF=Homo sapiens cDNA FLJ13303 fs, clone OVARC1001372, highly similar to Homo sapiens liprin-alpha4 mRNA. /FEA=mRNA /C	AK023365	1.02
Melanoma associated gene	BF342851	1.02
hypothetical protein FLJ23138	AW000928	1.02
teratocarcinoma repeat domain 3	AW510696	1.02
hypothetical protein MGC3282	NM_024029	1.02
erythroblast membrane-associated protein	NM_018538	1.02
ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)	BC005980	1.02
KH-type splicing regulatory protein (FUSE binding protein 2)	AI933301	1.02
paired box gene 3 (Waaerdenburg syndrome 1)	NM_013942	1.02
KIAA0532 protein	AI052003	1.02
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G	NM_017893	1.02
purinergic receptor P2X, ligand-gated ion channel, 3	NM_002559	1.02
Bardet-Biedl syndrome 4	AI813772	1.02
KIAA0401 protein	AB007861	1.02
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson	NM_022975	1.02
DKFZp566D133 protein	BC436400	1.02
hypothetical protein FLJ20871 similar to FSP27	NM_022094	1.02
neurogenic differentiation 8	NM_022728	1.02
pregnancy specific beta-1-glycoprotein 3	BC005824	1.02
phosphatidylinositol binding clathrin assembly protein	AL135735	1.02
solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2	AF55105	1.02
zinc finger protein 48 (KUP)	NM_008977	1.02
Consensus includes gb:AF052119.1 /DEF=Homo sapiens clone 23622 mRNA sequence /FEA=mRNA /DB_XREF=gi:3360428 /UG=Hs.151608 Homo sapiens clone 23622	AF052119	1.02
purinergic receptor P2X-like 1, orphan receptor	NM_005446	1.02
VIR-like 1	NM_020633	1.02
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2	NM_006854	1.02
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson	M87771	1.02
5' nucleoidase (CD73)	NM_002526	1.02
DNA (cytosine-5)-methyltransferase 2	AJ223333	1.02
bagpope homeobox homolog 1 (Drosophila)	NM_001189	1.02
Consensus includes gb:AK022254.1 /DEF=Homo sapiens cDNA FLJ12192 fs, clone MAMMA1000851. /FEA=mRNA /DB_XREF=gi:10433612 /UG=Hs.288451 Homo sapie	AK022254	1.02
aspartate beta-hydroxylase	AK021929	1.02
Consensus includes gb:AF049387.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N1918 (from clone DKFZp586N1918) /FEA=mRNA /DB_XREF=gi:4500178 /UG=Hs.16	AF049387	1.02
LIM domain only 7	NM_005358	1.02
Consensus includes gb:BF431965 /FEA=EST /DB_XREF=gi:11444079 /DB_XREF=est:na83609.x1 /CLONE=IMAGE:3274409 /UG=Hs.153106 Homo sapiens clone 2372	BF431965	1.02
casepase 10, apoptosis-related cysteine protease	AF111345	1.02
carbohydrate (N-acetyl)galactosamine 4-0) sulfotransferase 6	NM_022467	1.02
ten-1 homolog b (C. elegans)	NM_015322	1.02
glucose-6-phosphatase, transport (glucose-6-phosphate) protein 1	NM_001487	1.02
zinc finger, imprinted 2	NM_015383	1.02
catenin (cadherin-associated protein), delta 1	AB023382	1.02
phosphoinositide-3-kinase, regulatory subunit, polypeptide p101	NM_014308	1.02
retinoblastoma binding protein 5	NM_005057	1.02
bone morphogenetic protein 8 (osteogenic protein 2)	NM_001720	1.02
dermatopontin	NM_001937	1.02
ankyrin-like with transmembrane domains 1	NM_007332	1.02
glucose-6-phosphatase, catalytic (glycogen storage disease type I, von Gierke disease)	NM_000151	1.02
retinitis pigmentosa 2 (X-linked recessive)	NM_006915	1.02
lymphoid nuclear protein related to Ar-4	AF60465	1.02
tumor necrosis factor receptor superfamily, member 5	BF664114	1.02
ring finger protein 13	NM_007282	1.02
hypothetical protein FLJ20421	NM_017813	1.02
ATPase, Na+/K+ transporting, alpha 2 (*) polypeptide	NM_024718	1.02
protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I)	NM_000702	1.02
farnesyltransferase, CAAX box, beta	AL544951	1.02
	NM_002028	1.02

KIAA0781 protein
protein O-lucosyltransferase 1
Cluster incl. AL120741:DKFZp782K231_r1 Homo sapiens cDNA, 5' end /clone=DKFZp782K231 /clone_end=5 /gb=AL120741 /gi=5926641 /ug=Hs.8859 /len=707
eta homologous factor
neutrolisin
Consensus includes gb:AC005814 /DEF=Homo sapiens chromosome 19, cosmid F23289 /FEA=CDS /DB_XREF=gi:3540175 /UG=Hs.287391 Homo sapiens chromosome
Consensus includes gb:AF090100.1 /DEF=Homo sapiens clone IMAGE 23915 /FEA=mRNA /DB_XREF=gi:4063635 /UG=Hs.170241 Homo sapiens clone IMAGE 23915
KIAA1097 protein
chromosome 11 open reading frame 17
hypothetical protein
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13
p21/Cdc42/Rac1-activated kinase 1 (STE20 homolog, yeast)
SH3-domain binding protein 2
gb:NM_023939.1 /DEF=Homo sapiens hypothetical protein MGC2752 (MGC2752), mRNA, /FEA=mRNA /GEN=MGC2752 /PROD=hypothetical protein MGC2752 /DB_XREF=NM_023939
tumor necrosis factor (ligand) superfamily, member 15
Consensus includes gb:AI652058 /FEA=EST /DB_XREF=gi:4736037 /DB_XREF=gi:4736037 /DB_XREF=IMAGE:2306901 /UG=Hs.129081 Human Chromosome 16 BAC
KIAA0164 gene product
histone deacetylase 2
ret finger protein
glycerol kinase
KIAA0184 protein
PPAR binding protein
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 3
DKFZP568E104 protein
intersectin 1 (SH3 domain protein)
SPTF-associated factor 65 gamma
RAR-related orphan receptor A
ubiquitin specific protease 10
Consensus includes gb:AF035281.1 /DEF=Homo sapiens clone 23903 mRNA sequence, /FEA=mRNA /DB_XREF=gi:2661032 /UG=Hs.12526 Homo sapiens clone 23903
SELENOPHOSPHATE SYNTHETASE 1
selenium binding protein 1
hypothetical protein FLJ20897
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 3
hypothetical protein FLJ10890
pregnancy specific beta-1-glycoprotein 2
chromosome 12 open reading frame 3
ferritin, light polypeptide
ATP-binding cassette, sub-family C (CFTR/MRP), member 6
phosphatidylinositol transfer protein, membrane-associated
high-glucose-regulated protein 8
growth associated protein 43
sash-repeat protein
HIV TAT specific factor 1
purine-rich element binding protein A
KIAA0359 protein
CREB binding protein (Rubinstein-Taybi syndrome)
plasminogen activator, urokinase receptor
neutral sphingomyelinase
tumor necrosis factor receptor superfamily, member 6b, decoy
4-aminobutyrate aminotransferase
WD repeat domain 10
adenylate cyclase 3
lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)
connexin 59
hypothetical protein FLJ14360
platelet-derived growth factor receptor, alpha polypeptide
ncam
glutathione peroxidase 5 (epididymal androgen-related protein)
Snf2-related CBP activator protein
glycogenin 2
hypothetical protein FLJ13615
I-box and leucine-rich repeat protein 11
stromal antigen 1

AB018324 1.02
BC000582 1.02
AL120741 1.02
NM_012153 1.02
NM_006183 1.02
AC005814 1.02
AF090100 1.02
AK02864 1.02
NM_020642 1.02
NM_020313 1.02
AJ001698 1.02
U51120 1.02
AB000463 1.02
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NM_005118 1.02
NM_014739 1.02
NM_001527 1.02
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AC006293 1.02
AL110238 1.02
AF064243 1.02
AU154740 1.02
L14611 1.02
BG390445 1.02
AF035281 1.02
AV682679 1.02
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NM_004529 1.02
NM_018259 1.02
NM_031246 1.02
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NM_004910 1.02
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NM_014467 1.02
NM_014500 1.02
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NM_015113 1.02
NM_004380 1.02
U06839 1.02
NM_016317 1.02
NM_016434 1.02
NM_000663 1.02
NM_003996 1.02
AK023808 1.02
U94363 1.02
AF317887 1.02
BE675843 1.02
AW169948 1.02

acyl-Coenzyme A oxidase 1, palmitoyl
hypothetical protein FLJ20130
Consensus includes gb:AA534504 /FEA=EST /DB_XREF=gi:2271875 /DB_XREF=IMAGE:926246 /UG=Hs.8982 ESTs, Highly similar to KIAA1395
Janus kinase 1 (a protein tyrosine kinase)
cytochrome b-561
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1
FAST kinase
cyclin-dependent kinase inhibitor 1C (p57, Kip2)
interleukin 15
kinase suppressor of ras
heparanase
collagen, type IV, alpha 1
KIAA0653 protein, B7-like protein
Consensus includes gb:AB97981 /FEA=EST /DB_XREF=gi:5764799 /DB_XREF=est:wu1105.x1 /CLONE=IMAGE:2516673 /UG=Hs.115065 ESTs
chromosome 22 open reading frame 5
solute carrier family 4, sodium bicarbonate cotransporter, member 7
hypothetical protein FLJ11467
Consensus includes gb:AA769006 /FEA=EST /DB_XREF=gi:2820244 /DB_XREF=IMAGE:1318578 /UG=Hs.290821 ESTs, Highly similar to G027
heparan sulfate proteoglycan 2 (perlecan)
peptidyl-prolyl isomerase G (cyclophilin G)
cholecystokinin
msh homeo box homolog 2 (Drosophila)
Consensus includes gb:AK025422.1 /DEF=Homo sapiens cDNA: FLJ21769 fls, clone COL.F7354. /FEA=mRNA /DB_XREF=gi:10437930 /UG=Hs.306810 Homo sapiens cDNA: AK025422
hypothetical protein FLJ12871
pleomorphic adenoma gene-like 2
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 4
Janus kinase 3 (a protein tyrosine kinase, leukocyte)
protein tyrosine phosphatase, non-receptor type 7
hypothetical protein FLJ22527
erythropoietin receptor
interferon (alpha, beta and omega) receptor 2
Consensus includes gb:AF070538.1 /DEF=Homo sapiens clone 24420 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387895 /UG=Hs.159408 Homo sapiens clone 24420 mRNA sequence: AF070538
alpha2,3-sialyltransferase
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)
hyperpolarization activated cyclic nucleotide-gated potassium channel 2
Consensus includes gb:AK026682.1 /DEF=Homo sapiens cDNA: FLJ23029 fls, clone LNC01883. /FEA=mRNA /DB_XREF=gi:10439598 /UG=Hs.308864 Homo sapiens cDNA: AK026682
hypothetical gene DKFZp434A1010
endothelial differentiation, sphingolipid G-protein-coupled receptor, 5
lysophospholipase 1
chromosome 21 open reading frame 8
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 1
Consensus includes gb:AV647366 /FEA=EST /DB_XREF=gi:9868380 /DB_XREF=est:AV647366 /CLONE=GLCAXE09 /UG=Hs.282365 ESTs
solute carrier family 30 (zinc transporter), member 3
adenylyl cyclase-associated protein
Consensus includes gb:U82303.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938329 /UG=Hs.123080 Homo sapiens clone 24420 mRNA sequence: U82303
presenilin 2 (Alzheimer disease 4)
phytanoyl-CoA hydroxylase (Refsum disease)
hypothetical protein from EUROIMAGE 2021883
Cluster Inc. A1198543.q4911.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:1753389 /clone_end=3 /gb=A1198543 /gi=3751149 /ug=Hs.8982 /len=794
alpha-actinin-2-associated LIM protein
DNA binding protein for surfactant protein B
KIAA1085 protein
discs, large (Drosophila) homolog 1
trypartite motif-containing 9
myosin VI
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 4
tumor necrosis factor receptor superfamily, member 6b, decoy
interferon-stimulated protein, 15 kDa
KIAA0507 protein
transcobalamin I (vitamin B12 binding protein, R binder family)
pleiomorphic adenoma gene-like 1
sperm associated antigen 11
neurofibromin 2 (bilateral acoustic neuroma)

Consensus includes gb:AU147599 /FEA=EST /DB_XREF=gi:1009120 /DB_XREF=est:AU147599 /CLONE=MAMMA1001154 /UG=Hs.287501 Homo sapiens cDNA FLJ12
 galactose-4-epimerase, UDP- NM_000403 1.02
 tumor necrosis factor receptor superfamily, member 10b AF153887 1.02
 zinc finger protein NM_018651 1.02
 neuropathy target esterase NM_006702 1.02
 solute carrier family 16 (monocarboxylic acid transporters), member 3 AA527578 1.02
 myelin transcription factor 1 AB020842 1.02
 chymotrypsin-like AD72212 1.02
 ubiquitin specific protease 3 AF077040 1.02
 keratin, hair, basic, 1 NM_002281 1.02
 Sec23-interacting protein p125 AK001135 1.02
 interleukin 5 receptor, alpha M75914 1.02
 retinoic acid receptor responder (tazarotene induced) 1 NM_002888 1.02
 transmembrane 4 superfamily member 5 NM_003963 1.02
 nuclear LIM Interactor-interacting factor NM_021198 1.02
 ETAA18 protein NM_019002 1.02
 Consensus includes gb:AI992251 /FEA=EST /DB_XREF=gi:5839156 /DB_XREF=est:ws41d12.x1 /CLONE=IMAGE:2499767 /UG=Hs.184581 ESTs
 glycosylphosphatidylinositol specific phospholipase D1 NM_001503 1.02
 Consensus includes gb:BF448703 /FEA=EST /DB_XREF=gi:11514872 /DB_XREF=est:7n93902.x1 /CLONE=IMAGE:3572019 /UG=Hs.59728 Homo sapiens mRNA; cDNA BF448703 1.02
 crystallin, beta A1 NM_005208 1.02
 KIAA0763 gene product AB52645 1.02
 mucin 3A, intestinal AF007194 1.02
 Consensus includes gb:AA595276 /FEA=EST /DB_XREF=gi:2410626 /DB_XREF=est:tno35a05.s1 /CLONE=IMAGE:1102640 /UG=Hs.243804 Homo sapiens cDNA FLJ136 1.02
 v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian) AK024281 1.02
 RAB3B, member RAS oncogene family NM_002867 1.02
 natural killer-tumor recognition sequence AI361805 1.02
 GATA binding protein 4 AV700724 1.02
 DKFZP434J154 protein AK024279 1.02
 MAD, mothers against decapentaplegic homolog 6 (Drosophila) NM_005585 1.02
 growth differentiation factor 3 NM_020634 1.02
 ribosomal protein S12 BC002550 1.02
 AT2 receptor-interacting protein 1 AB95017 1.02
 KIAA0692 protein AB24382 1.02
 hypothetical protein DKFZp564B1162 NM_031305 1.02
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 7 AF212842 1.02
 Consensus includes gb:AW450910 /FEA=EST /DB_XREF=gi:6991886 /DB_XREF=est:UH-H-B13-alle-01-Q-UI.s1 /CLONE=IMAGE:2737225 /UG=Hs.74101 spleen tyrosine 1.02
 gb:LO8961.1 /DEF=Homo sapiens transmembrane tyrosine kinase mRNA, complete cds. /FEA=CDS /PROD=tyrosine kinase /DB_XREF=gi:697614 /FL-gb:LO8961.1 1.02
 neuro-oncological ventral antigen 2 L08961 1.02
 Consensus includes gb:R48779 /FEA=EST /DB_XREF=gi:810805 /DB_XREF=est:ty69c07.s1 /CLONE=IMAGE:153996 /UG=Hs.293845 ESTs, Weakly similar to B32688 bt R48779 1.02
 candidate tumor suppressor p33 ING1 homolog AA887063 1.02
 RYK receptor-like tyrosine kinase BG032035 1.02
 runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene) AF312387 1.02
 sialophorin (gp115, leukostallin, CD43) J04168 1.02
 PDZ domain containing 1 NM_002614 1.02
 capping protein (actin filament) muscle Z-line, beta U03271 1.02
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 6 BC004948 1.02
 tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein) AI275938 1.02
 microtubule-associated protein, RP/EB family, member 2 BE671156 1.02
 claudin 3 NM_001306 1.02
 renin binding protein NM_002910 1.02
 HIV-1 Rev binding protein AB98512 1.02
 natriuretic receptor 2 (cardiac) NM_001035 1.02
 Consensus includes gb:BF939092 /FEA=EST /DB_XREF=gi:12356412 /DB_XREF=est:705c01.x1 /CLONE=IMAGE:3706753 /UG=Hs.136090 ESTs 1.02
 alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I; glycolicaciduria; serine-pyruvate aminotransferase) BF939092 1.02
 prostaglandin E receptor 1 (subtype EP1), 42kD D13368 1.02
 hypothetical protein PRO0756 NM_000855 1.02
 calcium binding protein 5 NM_018583 1.02
 hypothetical protein NM_018855 1.02
 old astrocyte specifically induced substance Z62180 1.02
 semaphorin 5A AF055009 1.02
 sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A NM_003966 1.02
 aldolase B, fructose-bisphosphate BF195998 1.02
 LIM domain protein NM_003687 1.02
 apoptosis inhibitor 5 AF229254 1.02

ribosomal protein L31
 hypothetical protein FLJ13171
 solute carrier family 21 (organic anion transporter), member 9
 cytochrome P450, subfamily XXVIA, polypeptide 1
 protein phosphatase 1, regulatory (inhibitor) subunit 8
 Consensus includes gb:AL049351.1 /DEF=Homo sapiens mRNA: cDNA DKFZ566C093 (from clone DKFZ566C093). /FEA=mRNA /DB_XREF=gi:4500138 /UG=Hs.3020
 erythrocyte protein band 4.1-like 4
 Golgi apparatus protein 1
 small nuclear RNA activating complex, polypeptide 5, 19KD
 DKFZP588M1523 protein
 phosphoinositide-3-kinase, catalytic, gamma polypeptide
 nuclear mitotic apparatus protein 1
 IGF-II mRNA-binding protein 3
 poliovirus receptor
 KIAA0753 gene product
 hemochromatosis
 hypothetical protein
 likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K
 Consensus includes gb:BG420747 /FEA=EST /DB_XREF=gi:13327253 /DB_XREF=est:602448537 /1 /CLONE=IMAGE:4586984 /UG=Hs.259677 ESTs, Weakly similar to : BG420747
 anti-Müllerian hormone receptor, type II
 AND-1 protein
 palmitoyl-protein thioesterase 2
 KIAA0719 protein
 engulfment and cell motility 2 (ced-12 homolog, C. elegans)
 centrin, EF-hand protein, 2
 cytochrome c-like antigen
 thromboxane A2 receptor
 clathrin, light polypeptide (Lcb)
 heterogeneous nuclear ribonucleoprotein D-like
 paraxonase 3
 glypican 4
 methyl-CpG binding domain protein 2
 dyskeratosis congenita 1, dystekin
 keratin 6B
 nuclear protein 3 (apoptosis repressor with CARD domain)
 glucosidase, beta: add (includes glucosylceramidase)
 chloride intracellular channel 5
 natural killer-tumor recognition sequence
 gollioth protein
 PAN binding protein 9
 tanatodictin (beta)-like 1
 Cluster Incl. A1744451.wf89d08.x1 Homo sapiens cDNA, 3' end /clones=IMAGE-2362787 /clone_end=3 /gb=A1744451 /gi=5112739 /ug=Hs.124820 /len=584
 protein predicted by clone 23733
 low density lipoprotein receptor-related protein 3
 microfilament-associated protein 3
 KIAA0630 protein
 fibroblast growth factor receptor 4
 RAB5A, member RAS oncogene family
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2
 CREB-induced protein
 hypothetical protein FLJ12781
 adaptor-related protein complex 4, mu 1 subunit
 amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2)
 exosomes (multiple) 2
 putative dimethyladenosine transferase
 hypothetical protein FLJ20725
 Consensus includes gb:AF052100.1 /DEF=Homo sapiens clone 23645 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360407 /UG=Hs.6651 Homo sapiens clone 23645 m
 testis intracellular mediator protein
 Jun B proto-oncogene
 growth hormone 2
 serine/threonine kinase 13 (aurora/PL1-like)
 coagulation factor V (procoagulin, labile factor)
 TdT binding protein

disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)
protein disulfide isomerase, pancreatic
chromosome 11 open reading frame 24
cysteine and glycine-rich protein 2
cGMP deaminase
inositol 1,4,5-trisphosphate receptor, type 2
Consensus includes gb:BF19707 /FEA=EST /DB_XREF=gi:11087054 /DB_XREF=est:7m8n03.x1 /CLONE=IMAGE:3562997 /UG=Hs.155485 huntingtin interacting protein BF19707
Consensus includes gb:AL523184 /FEA=EST /DB_XREF=gi:12786677 /DB_XREF=est:AL523184 /CLONE=CSDDC001Y02 (5 prime) /UG=Hs.288146 ESTs, Highly similar
ERO1-like (S. cerevisiae)
calcium channel, voltage-dependent, alpha 1G subunit
son of sevenless homolog 1 (Drosophila)
FXRD domain-containing ion transport regulator 6
up-regulated by BCG-CWS
DKFZP586N1922 protein
PTK2 protein tyrosine kinase 2
cytochrome c oxidase subunit VIIb
Alport syndrome, mental retardation, midface hypoplasia and ellipocytosis chromosomal region, gene 1
KIAA0226 gene product
Consensus includes gb:AF070642.1 /DEF=Homo sapiens clone 24488 mRNA sequence. /FEA=mrRNA /DB_XREF=gi:3283915 /UG=Hs.323815 Homo sapiens clone 24488
RNA binding motif protein 8A
Consensus includes gb:AA664011 /FEA=EST /DB_XREF=gi:2618002 /DB_XREF=est:ac0310.s1 /CLONE=IMAGE:855403 /UG=Hs.173638 transcription factor 7-like 2 (T-) /AA664011
KIAA0603 gene product
Consensus includes gb:AL356115 /DEF=Human DNA sequence from clone RP11-488022 on chromosome 10 Contains the 3part of a gene for KIAA1128 protein, a novel
COQ7 coenzyme Q, 7 homolog ubiquinone (yeast)
KIAA0769 gene product
solute carrier family 4, anion exchanger, member 1 (erythrocyte membrane protein band 3, Diego blood group)
DnaJ (Hsp40) homolog, subfamily B, member 2
polymerase (RNA) II (DNA directed) polypeptide C (33KD)
hypothetical protein FLJ21324
homo box C10
rhabdoid tumor deletion region gene 1
neural cell adhesion molecule 1
adenosine monophosphate deaminase (isoform E)
Consensus includes gb:BE503392 /FEA=EST /DB_XREF=gi:9705800 /DB_XREF=est:7a17604.x1 /CLONE=IMAGE:3219006 /UG=Hs.145020 ESTs, Weakly similar to KIAA
hypothetical protein MGC10471
Ras association (RalGDS/AF-6) domain family 2
ATPase, Class V, type 10B
G1 to S phase transition 1
phospholipase C, gamma 1 (formerly subtype 14b)
6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 3
zinc finger protein 189
KIAA0941 protein
disrupted in schizophrenia 1
Consensus includes gb:AF028387 /FEA=EST /DB_XREF=gi:5664351 /DB_XREF=est:wc98f10.x1 /CLONE=IMAGE:2463211 /UG=Hs.331601 Homo sapiens, Similar to cyst
hypothetical protein
Consensus includes gb:AF070574.1 /DEF=Homo sapiens clone 24819 mRNA sequence. /FEA=mrRNA /DB_XREF=gi:3387945 /UG=Hs.22316 Homo sapiens clone 24819 /AF070574
proteasome (prosome, macropain) inhibitor subunit 1 (PI31)
Duffy blood group
Consensus includes gb:N57781 /FEA=EST /DB_XREF=gi:1201671 /DB_XREF=est:y56g05.s1 /CLONE=IMAGE:246776 /UG=Hs.269584 ESTs, Weakly similar to ALU5_H
UDP glycosyltransferase 8 (UDP-galactose ceramide galactosyltransferase)
immunoglobulin lambda joining 3
Consensus includes gb:AL139318 /DEF=Human DNA sequence from clone RP11-124817 on chromosome 13. Contains the DCT gene for dopachrome tautomerase (dopa
UDP glycosyltransferase 2 family, polypeptide B10
polymetric immunoglobulin receptor
hypothetical protein FLJ21174
tumor necrosis factor (TNF superfamily, member 2)
homeodomain interacting protein kinase 3
KIAA0403 protein
Consensus includes gb:AV694312 /FEA=EST /DB_XREF=gi:10296175 /DB_XREF=est:AV694312 /CLONE=GKCELG01 /UG=Hs.250638 Homo sapiens mRNA full length
arsenate resistance protein ARS2
carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7
protein phosphatase 1D magnesium-dependent, delta isoform

F-box only protein 3
neuron-specific protein
hypotheical protein FLJ13204
transcription factor 12 (HTF4, helix-loop-helix transcription factors 4)
endoplasmic reticulum oxidoreductin 1-Lbeta
KIAA0038 gene product
cytochrome P450, subfamily XXVIII (steroid 27-hydroxylase, cerebrotendinous xanthomatosis), polypeptide 1
N-myristyltransferase 2
adrenemic, alpha-2C-, receptor
RA838, member PAS oncogene family
hypotheical protein FLJ12401
PK508 binding protein 3 (25KD)
Consensus includes gb:AL031602 /DEF=Human DNA sequence from clone RP5-1174N9 on chromosome 1p34.1-35.3. Contains the gene for a novel protein with IBR dom
lysine 3-monoxygenase/hypophan 5-monoxygenase activation protein, theta polypeptide
KIT ligand
POU domain, class 3, transcription factor 4
Consensus includes gb:AU144900 /FEA=EST /DB_XREF=gi:11006421 /DB_XREF=est:AU144900 /CLONE=HEMBA1003327 /UG=Hs.274233 Homo sapiens cDNA FLJ10
T-cell leukemia/lymphoma 6
KIAA0774 protein
tRNA selenocysteine associated protein
growth factor receptor-bound protein 7
hypotheical protein MGC2716
Consensus includes gb:AA205593 /FEA=EST /DB_XREF=gi:1803801 /DB_XREF=est:z6806.s1 /CLONE=IMAGE:646782 /UG=Hs.26837 Homo sapiens mRNA; cDNA DI
melanoma cell adhesion molecule
hypotheical protein FLJ10408
wingless-type MMTV integration site family, member 5B
NIMA (never in mitosis gene a)-related kinase 1
hypotheical protein FLJ22794
fibulin 1
G protein-coupled receptor kinase 5
major histocompatibility complex, class II, DO alpha
KIAA1109 protein
olfactory receptor, family 1, subfamily F, member 1
nilogen-activated protein kinase kinase kinase 10
hypotheical protein MGC2550
toll-like receptor 3
KIAA0993 protein
CD5 antigen (p58-42)
MAD, mothers against decapentaplegic homolog (Drosophila) interacting protein, receptor activation anchor
actin-like 7B
hypotheical protein DKFZ586E1923
patched homolog (Drosophila)
Consensus includes gb:BE967207 /FEA=EST /DB_XREF=gi:11773627 /DB_XREF=est:601861094R1 /CLONE=IMAGE:3916174 /UG=Hs.165580 ribosomal protein S13
nuclear receptor subfamily 2, group F, member 6
thimet oligopeptidase 1
A kinase (PRKA) anchor protein (ytlaao) 9
CGI-48 protein
hypotheical protein MGC4845
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9
likely ortholog of preimplantation protein 3
methyl-CpG binding domain protein 2
lectin, mannose-binding, 1
Consensus includes gb:AI869947 /FEA=EST /DB_XREF=gi:4834721 /DB_XREF=est:w868d12.x1 /CLONE=IMAGE:2312567 /UG=Hs.71969 Homo sapiens mRNA; cDNA [AI869947
protein phosphatase 1, regulatory (inhibitor) subunit 3A (glycogen and sarcoplasmic reticulum binding subunit, skeletal muscle)
RNA, U transporter 1
hypotheical protein FLJ10525
KIAA0783 gene product
KIAA0609 protein
inter-alpha (globulin) inhibitor H4 (plasmin kallikrein-sensitive glycoprotein)
gb:BC001279.1 DEF=Homo sapiens. Similar to KIA0626 gene product, clone MGC:5129. mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0626 gene product
transducin-like enhancer of split 2 (Elsp1) homolog, (Drosophila)
Gem-interacting protein
ATP-binding cassette, sub-family C (CFTR/MRP), member 1

sec22 homolog
 CD8 antigen, beta polypeptide 1 (p37)
 5,10-methylenetetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase)
 basic leucine zipper transcription factor, ATF-like
 major histocompatibility complex, class I-like sequence
 lipophilin A (ulceroglobulin family member)
 natural killer cell group 7 sequence
 alpha-2-glycoprotein 1, zinc
 nuclear phosphoprotein similar to S. cerevisiae PWP1
 Rheus blood group-associated glycoprotein
 annexin A1
 voltage-dependent anion channel 1 pseudogene
 hypothetical protein MGC17330
 hypothetical protein
 thiamin pyrophosphokinase 1
 integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
 TAF_{II}116 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31 kD
 MLL, septin-like fusion
 Consensus includes gb:BG151527/FEA=EST/DB_XREF=gi:12633557/DB_XREF=est:nac28d07.x1/CLONE=IMAGE:3394597/UG=Hs.128055 ESTs, Moderately similar
 solute carrier family 7 (cationic amino acid transporter, y⁺ system), member 6
 guanine nucleotide binding protein (G protein), q polypeptide
 Ctip1-interacting zinc finger protein
 Recllin 1
 protein phosphatase 1, regulatory subunit 10
 solute carrier family 7 (cationic amino acid transporter, y⁺ system), member 6
 methylcrotonoyl-Coenzyme A carboxylase 1 (alpha)
 ring finger protein 15
 Consensus includes gb:A693140/FEA=EST/DB_XREF=gi:4970480/DB_XREF=est:wd66a05.x1/CLONE=IMAGE:2336720/UG=Hs.22907 Homo sapiens mRNA; cDNA
 cofactor required for Sp1 transcriptional activation, subunit 9 (33kD)
 neuro-oncological ventral antigen 1
 GREB1 protein
 alpha-methylacyl-CoA racemase
 BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)
 T. brachyury homolog (mouse)
 hypothetical protein P15-2
 hypothetical protein
 neural proliferation, differentiation and control, 1
 hypothetical protein FLJ10615
 hypothetical protein DKFZp434G131
 HCR (e-helix coiled-coil rod homologue)
 penitoxin-related gene, rapidly induced by IL-1 beta
 KIAA0471 gene product
 hypothetical protein DKFZp547D104
 cellular repressor of E1A-stimulated genes
 a disintegrin and metalloproteinase domain 15 (metagidrin)
 apical cord-derived growth factor-B
 hypothetical protein FLJ11286
 hypothetical protein FLJ20641
 hypothetical protein FLJ12895
 glial fibrillary acidic protein
 bromodomain adjacent to zinc finger domain, 1B
 ELL-RELATED RNA POLYMERASE II, ELONGATION FACTOR
 G protein-coupled receptor 45
 Consensus includes gb:AK025313.1 IDEF-Homo sapiens cDNA: FLJ21660 fis, clone COL08818, /FEA=mRNA/DB_XREF=gi:10437803/UG=Hs.287678 Homo sapiens cDNA
 solute carrier family 2, (facilitated glucose transporter) member 8
 interferon-stimulated transcription factor 3, gamma (48kD)
 Janus kinase 3 (e protein tyrosine kinase, leukocyte)
 hypothetical protein FLJ13725
 zona pellucida binding protein
 HSPC056 protein
 hypothetical protein FLJ10211
 preludin 4
 Cluster Ind. A174083:wc38b04.x1 Homo sapiens cDNA, 3 end clone=IMAGE:2317231/clone_end=3 /gb=A1744083 /gi=5112371 /ug=Hs.65406 /len=608

myosin VA (heavy polypeptide 12, myosin)	ZZ2957	1.01
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 38	AF038391	1.01
glucagon-like peptide 1 receptor	U01104	1.00
endothelin receptor type B	NM_003991	1.00
complement component 8, gamma polypeptide	M17253	1.00
ubiquitination factor E4B (UFD2 homolog, yeast)	NM_006048	1.00
stannin	NM_003498	1.00
CGI-60 protein	NM_016008	1.00
pyruvate dehydrogenase phosphatase	NM_018444	1.00
hypothetical protein DKFZp547D146	NM_020217	1.00
Consensus includes gb:AL358602.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547D146 (from clone DKFZp547D146) /FEA=mRNA /DB_XREF=gi:8655670 /UG=Hs.3065	AL358602	1.00
KIAA0545 protein	AB011117	1.00
glutathione S-transferase M1	X08020	1.00
G protein-coupled receptor 68	NM_006056	1.00
adenylate cyclase 1 (brain)	L05500	1.00
leucine-rich repeat protein, neuronal 3	AL442092	1.00
lary-RNA synthetase	NM_005548	1.00
hypothetical protein FLJ20532	BC420893	1.00
Consensus includes gb:A027701 /FEA=EST /DB_XREF=gi:5663665 /DB_XREF=est:wo5h11.x1 /CLONE=IMAGE:2462181 /UG=Hs.25307 Homo sapiens clone 24812 ml /A027701	BC420893	1.00
LIM and cysteine-rich domains 1	NM_014583	1.00
3-oxoacid CoA transferase	NM_000436	1.00
Consensus includes gb:AW969913 /FEA=EST /DB_XREF=gi:8159757 /DB_XREF=est:EST381991 /UG=Hs.105121 ESTs	AW969913	1.00
calbindin 3, (Vitamin D-dependent calcium binding protein)	NM_004057	1.00
Consensus includes gb:A095896 /FEA=EST /DB_XREF=gi:3434872 /DB_XREF=est:qb21a10.x1 /CLONE=IMAGE:1686890 /UG=Hs.301463 Human Chromosome 16 BAC	AO95896	1.00
N-ethylmaleimide-sensitive factor attachment protein, alpha	NM_003827	1.00
chromosome 11 open reading frame 15	NM_020644	1.00
synaptosomal-associated protein, 23kD	BC003686	1.00
AIM-1 protein	NM_016180	1.00
hypothetical protein MGC14817	AF255792	1.00
Consensus includes gb:AF19846.1 /DEF=Homo sapiens PRO1474 mRNA, complete cds. /FEA=mRNA /PROD=PRO1474 /DB_XREF=gi:7770128 /UG=Hs.324925 LATS1	AF19846	1.00
DiGeorge syndrome critical region gene 6	NM_006765	1.00
vascular Rab-GAP/TBC-containing	NM_007063	1.00
KIAA0663 gene product	AB032116	1.00
colipase, pancreatic	NM_001832	1.00
SMA5	X83301	1.00
lysosomal-associated membrane protein 1	A1248598	1.00
protophyll Coenzyme A carboxylase, alpha polypeptide	NM_000282	1.00
complex-associated-testis-expressed 1-like 1	NM_005519	1.00
DKFZP434J214 protein	AL556438	1.00
myosin IB	AK022489	1.00
CASP8 and FADD-like apoptosis regulator	AF005775	1.00
Rab geranylgeranyltransferase, beta subunit	AA128753	1.00
hydroxysteroid (17-beta) dehydrogenase 4	NM_000414	1.00
KIAA1088 protein	AB029009	1.00
ankyrin repeat and SOCS box-containing 1	AF055024	1.00
hypothetical protein MGC5178	NM_024044	1.00
zinc finger protein 134 (clone pHZ-15)	AC003682	1.00
G protein-coupled receptor, family C, group 5, member D	NM_018654	1.00
Consensus includes gb:X01695.1 /DEF=H.sapiens GENX-5624 mRNA, 3 UTR. /FEA=mRNA /DB_XREF=gi:2370158 /UG=Hs.194765 H.sapiens GENX-5624 mRNA, 3 UTR	AL080191	1.00
DKFZP434J082 protein	NM_000049	1.00
aspartoacylase (aminoadacylase 2, Canavan disease)	AW294107	1.00
succinate dehydrogenase complex, subunit B, iron sulfur (lp)	NM_005962	1.00
MAX interacting protein 1	NM_000287	1.00
peroxisomal biogenesis factor 6	NM_006382	1.00
chromosome 17 open reading frame 1A	NM_031244	1.00
shuin silent mating type information regulation 2 homolog 5 (S. cerevisiae)	AL536966	1.00
guanine nucleotide binding protein (G protein), gamma 3	AA523444	1.00
putative c-Myc-responsive	NM_000584	1.00
interleukin 8	NM_012463	1.00
TJ6 protein	NM_002821	1.00
PTX7 protein tyrosine kinase 7	NM_016061	1.00
CGI-127 protein	NM_001912	1.00
cathepsin L		

neuroglobin
PET112-like (yeast)
Consensus includes gp:X81637.1 /DEF=H.sapiens clathrin light chain b gene, /FEA=mrna, DB_XREF=gi:963046 /UG=Hs.73919 clathrin, light polypeptide (Lcb)
cytochrome P450, subfamily 11E (ethanol-inducible)
splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor)
KIAA0494 gene product
protocadherin 9
IQ motif containing GTPase activating protein 1
sigma receptor (SR31747 binding protein 1)
Consensus includes gp:AA906578 /FEA=EST, DB_XREF=est:02206.s1 /CLONE=IMAGE:1524202 /UG=Hs.23158 ESTs
solute carrier family 25 (mitochondrial carrier, peroxisomal membrane protein, 34kD), member 17
Consensus includes gp:AI967981 /FEA=EST, DB_XREF=gi:5764779 /DB_XREF=est:wu11008.x1 /CLONE=IMAGE:2516655 /UG=Hs.26837 Homo sapiens mRNA; cDNA [AI967981
phosphatidylinositol (4,5) biphosphate 5-phosphatase, A
ZYG homolog
hypothetical protein FLJ20224
glycan 1
NAD(P)H dehydrogenase, quinone 1
antigen identified by monoclonal antibody Ki-67
BCL2/adrenovirus E18 18kD interacting protein 3-like
RAV binding protein 9
hypothetical protein FLJ23393
nuclear transcription factor Y, alpha
vacuolar protein sorting 41 (yeast)
zinc finger protein, X-linked
calcium channel, voltage-dependent, alpha 1G subunit
hypothetical protein FLJ20039
integrin, beta 2 (antigen CD18 (p95), lymphocyte function-associated antigen 1; macrophage antigen 1 (mac-1) beta subunit)
homo box C4
anaplastic lymphoma kinase (K1-1)
hypothetical protein FLJ23416
plasminogen-like
jumonji homolog (mouse)
oligodendrocyte myelin glycoprotein
hypothetical protein FLJ11175
delta-like homolog (Drosophila)
neirin 1
beta-transducin repeat containing
ectodermal dysplasia 1, anhidrotic
KIAA1111 protein
hypothetical protein FLJ22060
leucine-rich repeat protein, neuronal 1
K01391 B subtilis TrpE protein, TrpD protein, TrpC protein corresponding to nucleotides 1883-4400 of K01391 [-S, -M, -3 represent transcript regions 5 prime, Middle, and 3 K01391
hypothetical protein R26124_1
DKFZP566H073 protein
fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Ptaiffer syndrome)
gp:NM_005993.1 /DEF=Homo sapiens T-box 10 (TBX10) mRNA, /FEA=mrna, /GEN=TBX10 /PROD=T-box 10 /DB_XREF=gi:5174708 /UG=Hs.203847 T-box 10 /FL=gp:NM_005993
Consensus includes gp:AU151793 /FEA=EST, DB_XREF=gi:11013314 /DB_XREF=est:AU151793 /CLONE=NT2RP2008115 /UG=Hs.157078 Homo sapiens cDNA FLJ127
hypothetical protein FLJ12409
macrophage receptor with collagenous structure
Consensus includes gp:AI207120 /FEA=EST, DB_XREF=gi:3765792 /DB_XREF=est:q59a09.x1 /CLONE=IMAGE:1754344 /UG=Hs.248129 glutathione peroxidase 5 (epid A207120
heal shock 105kD
Consensus includes gp:AW167793 /FEA=EST, DB_XREF=gi:63599401 /DB_XREF=est:q56007.x1 /CLONE=IMAGE:2632333 /UG=Hs.164038 Homo sapiens AKAP350C n AW167793
hypothetical protein FLJ20321
hypothetical protein MGCA309

NM_021257
NM_004564
X81637
AF162278
M69040
BC002525
AI524125
NM_003870
AF226804
AA906578
AL048784
AI967981
AI935720
NM_006336
NM_017719
NM_002081
BC000906
AU147044
AF060922
AA191426
NM_024639
AL031776
NM_014396
X59740
AB012043
NM_017635
NM_000211
AW298598
U66559
AK027069
NS6120
NM_004973
NM_002544
NM_018349
U15979
NM_004822
AF101784
AF081191
AW248934
NM_024612
AI631881
NM_024892
NM_021391
AC005626
AK025329
M80485
NM_005995
AU151793
NM_025105
NM_008770
AI207120
NM_006844
NM_017786
NM_024115

FIGURE 4C.

Gene	NCBI Accession Number	FOLD INCREASE
gp.L23516.1 /DEF=Human Ig rearranged gamma-chain, V-DJF+JHk, complete cds. /FEA=mRNA /DB_XREF=gi:385218 /FL=gb:L23516.1	L23516	43.37
ubiquitin carrier protein	NM_014501	41.83
glyceraldhyde-3-phosphate dehydrogenase	M33197	41.07
Lysosomal-associated multispansing membrane protein-5	NM_006762	37.81
ribosomal protein L29	NM_000992	37.44
gp.L23518.1 /DEF=Human Ig rearranged gamma-chain, V-DJF+JHk, complete cds. /FEA=mRNA /DB_XREF=gi:385220 /FL=gb:L23518.1	L23518	37.23
immunoglobulin heavy constant mu	U00139	33.58
adaptor-related protein complex 2, mu 1 subunit	NM_004068	32.97
heat shock 90D protein 1, beta	A218219	32.61
neurospage migration inhibitory factor (glycosylation-inhibiting factor)	NM_002415	30.90
lentin, light polypeptide	B538364	30.50
ornithine decarboxylase antizyme 1	A709094	30.15
Consensus includes gi:148784 /DEF=OSO Homo sapiens cDNA, FEA=mRNA /DB_XREF=gi:1066715 /UG=Hs.192426 ribosomal protein 32	L48784	30.09
endlease 1, (alpha)	U88968	28.76
ribosomal protein L29	BF683426	28.60
ribosomal protein L18	NM_000979	28.20
ribosomal protein L16	NM_000980	28.19
chaperone containing TDP1, subunit 7 (eta)	NM_006429	27.91
immunoglobulin heavy constant mu	NM_001536	25.89
MYT1 mitogen-activated protein kinase 2 (S. cerevisiae)	B3340548	27.05
protein 1	NM_005022	24.23
procollagen-proline, 2-oxoglutarate 4-lyase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase, thyroid hormone binding protein p55)	NM_000918	23.95
procollagen-proline, 2-oxoglutarate 4-lyase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase, thyroid hormone binding protein p55)	J02783	23.56
Consensus includes gb:084143.1 /DEF=Human immunoglobulin (mAb59) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V region /DB_XREF=084143	084143	22.30
Consensus includes gb:A249377.1 /DEF=Homo sapiens partial mRNA for human Ig lambda light chain variable region, clone MB91 (331 bp). /FEA=mRNA /GEN=IGL /PROD=immunoglobulin A249377	A249377	21.91
U6 snRNA-associated SnRNP protein L5m7	NM_016199	21.63
polymerase (DNA directed), delta 2, regulatory subunit (SNU2)	NM_006230	21.62
ribosomal protein L8	NM_000973	21.19
translocase (Nucleoside-korauril syndrome)	BF686840	21.11
gp.L235092.1 /DEF=Homo sapiens mRNA for single-chain antibody, complete cds. /FEA=cDS /PROD=immunoglobulin /DB_XREF=gi:3090425 /UG=Hs.249245 Homo sapiens mRNA for sin A2325092	A2325092	20.48
T-cell leukemia/lymphoma 1A	A22240	19.28
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	A910614	19.19
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	A910614	19.08
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	B000533	18.47
gp.L07950.1 /DEF=Homo sapiens HIC class 1 HLA B71 mRNA, complete cds. /FEA=cDS /GEN=HLA-B /PROD=HLA B71 /DB_XREF=gi:307236 /FL=gb:L07950.1	L07950	17.83
glyceraldhyde-3-phosphate dehydrogenase	M33197	17.71
ACOT7 mitochondrial acyl-CoA thioesterase 7 (S. cerevisiae)	A279900	17.26
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	NM_003751	16.97
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	NM_002654	16.81
pyruvate kinase, muscle	M87338	16.75
replication factor 5 (p53-binding protein 1) 2 (4202)	BF689355	16.61
glyceraldhyde-3-phosphate dehydrogenase	NM_003132	16.57
synovial synovial	NM_005466	16.27
RNA polymerase II transcriptional regulation mediator (Med6, S. cerevisiae, homolog of)	NM_006423	16.24
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	A910614	16.05
ribosomal protein L13	A910614	15.99
Consensus includes gb:084140.1 /DEF=Human immunoglobulin (mAb56) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V region /DB_XREF=084140	084140	15.89
CD22 antigen	552785	15.89
neutrophil cytosolic factor 4 (4202)	NM_013416	15.70
small EDRB-rich factor 2	NM_005770	15.52
glyceraldhyde-3-phosphate dehydrogenase	BE561479	15.36
translocase (Nucleoside-korauril syndrome)	L12711	15.06
ATP synthase, H ⁺ -transporting, mitochondrial F1 complex, delta subunit	BE798517	14.96
CSG-135 protein	NM_016068	14.94
glucose phosphate isomerase	NM_000175	14.88
peptidase, disulfide isomerase associated protein	A712222	14.86
actin, beta	J00351	14.30
Consensus includes gb:A254422 /DEF=Homo sapiens SHAR (A4) isoform (SHAR(A4)) gene, complete cds, alternatively spliced /FEA=cDS_2 /DB_XREF=gi:10946127 /UG=Hs.78202 SWISN A254422	A254422	14.20
lactin, derived from (11:2:16) malignant liposarcoma	NM_004960	14.10
ribosomal protein S5	NM_001009	13.79
polymerase (RNA) II (DNA directed) polypeptide 1 (13.3A0)	B6335629	13.39

heat shock 90kD protein 1, beta	AF275119	13.32
nuclease sensitive element binding protein 1	BE966374	13.25
ATP synthase, H ⁺ -transporting, mitochondrial F1 complex, delta subunit	NM_001687	13.24
euclaryotic translation initiation factor 3, subunit 8 (110kD)	AA679705	13.07
immunoglobulin heavy constant mu	BC001872	13.03
actin, beta	X00351	12.99
serine hydroxymethyltransferase 2 (mitochondrial)	NM_005412	12.93
polymerase (RNA) II (DNA directed) polypeptide E (25kD)	A554759	12.83
maternal 610 transcript	NM_003910	12.80
major histocompatibility complex, class I, B	083043	12.59
cell death-regulatory protein 6 (BIM15)	NM_015965	12.46
proteasome (prosome, macropain) 26S subunit, ATPase, 3	AL545823	12.22
velutin-containing protein	AF100752	12.15
nuclear RNA helicase, DEAD variant of DEAD box family	NM_005804	11.98
euclaryotic translation initiation factor 3, subunit 8 (110kD)	NM_003752	11.83
coronin, actin binding protein, 1A	U34690	11.81
mitochondrial ribosomal protein S12	NM_021107	11.76
cold inducible RNA binding protein	NM_001280	11.67
DEAD-box protein abstract	NM_016222	11.65
putative transmembrane protein; homolog of yeast Gcd3 membrane protein Yrip (Yip1p-interacting factor)	NM_020470	11.52
protein phosphatase 1c (formerly 2c), magnesium-dependent, gamma isoform	NM_002707	11.46
immunoglobulin heavy constant mu	574639	11.35
acid phosphatase 1, soluble	BC035989	11.32
G6a-51 protein	NM_015380	11.29
ATP synthase, H ⁺ -transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1	AL080089	11.11
7-dehydrocholesterol reductase	AW150953	11.06
B lymphoid tyrosine kinase	NM_001715	11.05
chaperonin containing TCP1, subunit 3 (gamma)	NM_005998	10.96
cell division cycle 2-like 2	AF067524	10.85
dehydro-diphosphoglycerate-kinase-protein glycosyltransferase	D25643	10.84
transforming growth factor, beta 1	BC000125	10.80
dephospho-cytochrome c reductase core protein 1	NM_003365	10.71
MDH1 dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (90D, B9)	NM_004542	10.61
mitochondrial ribosomal protein S2	NM_016034	10.58
chaperonin containing TCP1, subunit 2 (beta)	AL545982	10.54
ribosomal protein L18a	AC004692	10.40
polymerase (RNA) II (DNA directed) polypeptide 1 (13.3kD)	AW402635	10.24
interferon stimulated gene (20kD)	NM_002201	10.17
gp-M24668.1 (DEF=Human Ig V-nearby H-chain V-region mRNA (C-D-JH4), complete cds. /FEA=nmRNA/GEN=IGH/D8_XREF=gi:185198 /FL=gb-M24668.1	M24668	10.01
proteasome (prosome, macropain) 26S subunit, non-ATPase, 8	NM_002812	9.92
proteasome (prosome, macropain) subunit, alpha type, 7	AL078633	9.84
ribosomal protein S19	BE239729	9.82
major histocompatibility complex, class II, DR beta 5	AC297586	9.71
bresein 2 (60-70kD)	AA284075	9.67
similar to HYPOTHETICAL 34.0 XDA PROTEIN Z0795.3 IN CHROMOSOME IV	BE747342	9.65
phosphoserine aminotransferase	NM_021154	9.50
sukita carrier family 2 (facilitated glucose/fructose transporter), member 5	BE560461	9.50
mcd5 homolog 6 (E. coli)	D89646	9.48
gp-M24624.1 (DEF=Homo sapiens HLA class I heavy chain (HLA-Cw*1701) mRNA, complete cds. /FEA=CD5/GEN=HLA-Cw*1701 /PROD=HLA class I heavy chain /D8_XREF=gi:1575443 /FL=U02824	U02824	9.44
stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein)	L42024	9.43
nucleosome assembly protein 1-like 4	BE86580	9.39
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	NM_005969	9.31
ribosomal protein S19	A033605	9.29
replication factor C (activator 1) 2 (40kD)	NM_001022	9.29
euclaryotic translation initiation factor 4 gamma, 1	NM_002914	9.28
glutathione S-transferase pi	AF104913	9.27
signal sequence receptor, delta (translocin-associated protein delta)	NM_000852	9.15
VP528 protein	NM_016208	9.01
DMP7-564H182 protein	AK000022	8.99
6-phosphogluconolactonase	NM_012088	8.94
ribosomal protein S2	NM_002952	8.88
protein phosphatase 1, regulatory subunit 7	BE718769	8.85

ribosomal protein S15	NM_001018	8.78
translocase of inner mitochondrial membrane 13 homolog 8 (yeast)	NM_012458	8.77
endothelial differentiation-related factor 1	AB022582	8.75
T cell receptor beta locus	M15564	8.72
hypothetical protein R33729_1	AC005339	8.61
ubiquitin C	M26880	8.53
major histocompatibility complex, class II, DR alpha	M60333	8.52
non-FOU-domain-containing, octamer-binding	L14599	8.52
major histocompatibility complex, class II, DR alpha	M60334	8.36
DNA replication factor	AF321125	8.34
PRP8 pre-mRNA processing factor 8 homolog (yeast)	NM_006445	8.28
SBR1 homolog (S. pombe)	NM_006109	8.28
adrenyctic translation elongation factor 1 alpha 1	AL035687	8.27
KIA04217 protein	BC033381	8.24
major histocompatibility complex, class II, DR beta 1	U05585	8.20
ubiquitin-activating enzyme E1 (A15ST and BMT5 temperature sensitivity complementing)	NM_003334	8.14
polymerase (RNA) III (DNA directed) polypeptide K (12.3 kDa)	NM_016310	8.11
ATP synthase, H ⁺ -transporting, mitochondrial FO complex, subunit f, isoform 2	NM_004889	8.06
glutathionyl-HNA synthetase	NM_005051	8.04
HLA-B associated transcript 1	NM_004640	8.01
Consensus includes gp:AF043584.1 /DEF=Homo sapiens clone ASHmg1-b1 immunoglobulin lambda chain VJ region, (J6L) mRNA, partial cds. /FEA=mRNA /GEN=HL /PROD=immunoglobul	AF043584	8.00
ribosomal protein S14	AF116710	7.98
interferon regulatory factor 4	NM_002460	7.94
T cell receptor beta locus	AF043179	7.91
HLA-B associated transcript 3	BC028444	7.90
bone marrow stromal cell antigen 2	NM_004335	7.84
integrin beta 4 binding protein	AF022229	7.76
putative breast adenocarcinoma marker (32kb)	NM_014453	7.65
polymerase (RNA) II (DNA directed) polypeptide L (7.6kD)	BC005903	7.63
hypothetical protein PRO1847	AF119855	7.60
D123 gene product	NM_006023	7.54
actin binding LIM protein 1	NM_006720	7.53
adrenyctic translation initiation factor 3, subunit 2 (beta, 36kD)	U35764	7.47
T cell receptor beta locus	AL591122	7.45
tubulin, beta, 4	AL569749	7.39
customer protein complex, subunit epsilon	NM_007263	7.35
chromosome 14 open reading frame 3	NM_012111	7.35
major histocompatibility complex, class II, DP alpha 1	M27487	7.30
mitochondrial ribosomal protein S34	NM_023936	7.28
Hepatitis delta antigen-interacting protein A	NM_006848	7.27
pepdylprolyl isomerase F (cyclophilin F)	NM_005729	7.26
nuclear RNA export factor 1	BC004904	7.24
ubiquitin C	AB009010	7.21
transmembrane protein 4	BC001027	7.20
erythroid beta (AE) precursor-like protein 2	AW001847	7.19
hypothetical protein FL20113	AL523776	7.19
GM1 general control of amino-acid synthesis 1-tila 1 (yeast)	AB91055	7.16
MIF2 non-histone chromosome protein 2-tila 1 (S. cerevisiae)	AF155235	7.15
adrenyctic translation initiation factor 4A, isoform 1	NM_001416	7.13
macrophage erythroidblast attach	NM_005882	7.11
chemokine (C-X-C motif), receptor 4 (Itzin)	AF348491	7.10
calreticulin	AD000092	7.09
ancient ubiquitous protein 1	NM_012103	7.07
enoyl-CoA hydratase	AF0702405	7.06
putative human HLA class II associated protein 1	BE560202	7.05
suppression of tumorigenicity 14 (colon carcinoma, metastase, epithin)	NM_021978	7.05
proteasome (prosome, macropain) 26S subunit, non-ATPase, 3	NM_002809	7.02
GDP-mannose 4,6-dehydratase	NM_001500	7.01
transcription factor Dp-1	NM_007111	7.01
w-aryp myeloblastosis viral oncogene homolog (wim) -like 2	NM_002466	6.90
origin recognition complex, subunit 5-tila (yeast)	AF081459	6.89
Consensus includes gp:AF005447.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6-0201 allele, sequence. /FEA=mRNA /DB_XREF=gi:5515893 /LG=Hs.167385 Homo	AF005487	6.89
splicing factor, arginine/serine-rich 5	AW084582	6.88

Consensus includes g08E905165 (FEA=EST /DB_XREF=gi:177184 /DB_XREF=est:601166685T1 /CLONE=HMAE2959580 /UG=Hs. 100623 phospholipase C, beta 3, neighbor pseudogen	BE305165	6.88
ATP binding cassette, sub-family F (ABC2), member 2	NM_005692	6.87
adenoviral protein complex 2, sigma 1 subunit	NM_021575	6.86
U6 snRNA-associated Sm-like protein	NM_012321	6.86
unknown	NM_016209	6.85
major histocompatibility complex, class I, F	AW514210	6.84
ribosomal protein S2	A183766	6.82
H5V1 binding protein	NM_018694	6.81
CD79A antigen (immunoglobulin-associated alpha)	NM_001783	6.81
major histocompatibility complex, class II, DR beta 4	NM_021983	6.80
FTS66 binding protein precursor	NM_016594	6.77
FTS66 binding protein precursor	BC000733	6.77
eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD)	AI380004	6.76
ADP-ribosylation factor 1	NM_016459	6.74
hypothetical protein	BC001081	6.74
hypothetical protein	AI12507	6.69
anaphase promoting complex subunit 5	NM_001961	6.68
U6 snRNA-associated Sm-like protein	AL118502	6.67
eukaryotic translation elongation factor 2	BC068249	6.66
hypothetical protein M624675	NM_022804	6.64
guanylate kinase 1	A832239	6.62
SHRN upstream reading frame	BC003629	6.58
mitochondrial ribosomal protein L23	X12530	6.57
RNA, U2, small nuclear	NM_016292	6.51
membrane-spanning 4-domains, subfamily A, member 1	NM_003564	6.50
head shock protein 75	M61715	6.47
transglutinin 2	AL567820	6.46
trypophanyl-tRNA synthetase	BC066210	6.45
actin, gamma 1	A741124	6.45
eukaryotic translation initiation factor 4A, isoform 1	A953822	6.43
guanine nucleotide binding protein (G protein), beta polypeptide 1	A692974	6.42
ribosomal protein, large, P0	NM_002808	6.42
ribonucleotide reductase M1 polypeptide	NM_006427	6.38
proteasome (prosome, macropain) 26S subunit, non-ATPase, 2	J04564	6.36
CD27-binding (Siv) protein	BC002807	6.33
small nuclear ribonucleoprotein polypeptides B and B1	NM_001002	6.32
membrane-spanning 4-domains, subfamily A, member 1	BC005863	6.30
ribosomal protein, large, P0	AI109955	6.28
ribosomal protein, large, P0	BC003655	6.28
sub4D	NM_016065	6.26
ribosomal protein, large, P0	A503111	6.23
mitochondrial ribosomal protein S16	NM_004599	6.23
CD27-binding (Siv) protein	NM_001803	6.21
zinc regulatory element binding transcription factor 2	NM_006013	6.21
CDWS2 antigen (CAPIX11-1 antigen)	NM_001861	6.19
ribosomal protein L10	U14990	6.17
cytochrome c oxidase subunit IV isoform 1	A972144	6.15
ribosomal protein S3	NM_006631	6.15
hydroxymethyl-Coenzyme A dehydrogenase/3-oxoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	A792131	6.14
neurosp3 cytosolic factor 4 (40kD)	NM_001098	6.11
NADH dehydrogenase (ubiquinone) flavoprotein 1 (51kD)	NM_001029	6.11
acetylase 2, mitochondrial	AF226990	6.10
ribosomal protein S26	A744900	6.08
HLA-E histocompatibility antigen, class I, E	NM_006114	6.08
SWI/SNF related, actin associated, actin dependent regulator of chromatin, subfamily a, member 4	A438978	6.05
DNA segment on chromosome 19 (unique) 1177 expressed sequence	NM_005356	6.04
ribosomal protein L13	NM_002799	6.03
proteasome (prosome, macropain) subunit, beta type, 7	NM_005356	6.04
lymphocyte-specific protein tyrosine kinase	NM_006513	6.03
seyl-dRNA synthetase	M79321	6.02
v-pes-1 Yamaguchi sarcoma viral related oncogene homolog	BC006244	5.99
HSPK142 protein	BC395660	5.98
replication protein A1 (70kD)	86258784	5.96
ubiquitin-conjugating enzyme E26 2 (UBK7 homolog, yeast)		
membrane component, chromosome 11, surface marker 1		

H1 histone family, member X	NM_006026	5.95
lysosomal-associated multispinning membrane protein-5	A039086	5.95
PTD008 protein	NM_016145	5.95
poly(I:C) binding protein 1	U24223	5.94
polyglutamine binding protein 1	A041836	5.93
active homology 2 (Troxopala)	BC000422	5.93
MAOS box transcription enhancer factor 2, polypeptide 8 (myocyte enhancer factor 2B)	NM_005919	5.92
translocase of inner mitochondrial membrane 44 homolog (yeast)	NM_006351	5.92
intercellular adhesion molecule 2	NM_000873	5.91
bromodomain-containing 2	D42040	5.91
lysosomal-associated membrane protein 1	NM_005561	5.91
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, A5H1)	NM_005004	5.87
actin related protein 2/3 complex, subunit 2 (34 kD)	A273893	5.84
Consensus includes gb:83344495 (FJ451718) /gb_XREF=gi:13140933 /gb_XREF=est:6024611281 /CLONE=IMAGE-4377718 /UG=Hs.228201 ESTs, Moderately similar to AF118094.25 Pf	NM_004135	5.80
bicucitine dehydrogenase 3 (NMH+) gamma	NM_007108	5.80
mitochondrial ribosomal protein L9	A049636	5.80
transcription elongation factor 8 (SIII), polypeptide 2 (18kD, elongin B)	A670847	5.80
hypothetical protein	NM_014145	5.77
HSPC274 protein	M60261	5.77
APX1 nuclease (multifunctional DNA repair enzyme)	NM_016185	5.75
hematological and neurological expressed 1	NM_025205	5.75
hypothetical protein D872434H185	A047695	5.75
calcylcic translation initiation factor 4E-like 3	NM_001640	5.74
N-acylfluoroyl-peptide hydrolase	A452514	5.73
QMA0746 protein	A0001527	5.72
capitin, small subunit 1	AF147209	5.69
interleukin enhancer binding factor 3, 90kD	NM_006773	5.67
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	AL049650	5.67
hypothetical protein M610715	AF034607	5.66
chloride intracellular channel 1	NM_000626	5.64
CD79b antigen (immunoglobulin-associated beta)	NM_004493	5.64
hydroxymethyl-Coenzyme A dehydrogenase, type II	B1871375	5.62
zinc finger protein 207	AC058111	5.61
ribosomal protein S28	A028970	5.61
synactin 15	NM_012336	5.61
nuclear prelamin A recognition factor	NM_000101	5.60
cytochrome b-245, alpha polypeptide	NM_006409	5.60
actin related protein 2/3 complex, subunit 1A (41 kD)	NM_021126	5.57
metacryptopyruvate sulfotransferase	B1979419	5.57
Consensus includes gb:87979419 (FEA=EST) /gb_XREF=gi:12346634 /gb_XREF=est:602288246F1 /CLONE=IMAGE-4373914 /UG=Hs.119122, ribosomal protein L13a	J02523	5.57
lymphocyte cytosolic protein 1 (L-calcin)	NM_004551	5.56
NADH dehydrogenase (ubiquinone) Fe-S protein 3 (30kD) (NADH-coenzyme Q reductase)	NM_004596	5.55
small nuclear ribonucleoprotein polypeptide A	B342707	5.55
pre-mRNA processing factor 31 homolog (yeast)	NM_002708	5.54
protein phosphatase 1, catalytic subunit, alpha isoform	NM_003321	5.54
1a translation elongation factor, mitochondrial	A4514622	5.54
QMA0518 gene product	U34994	5.51
protein kinase, (DNA)-activated, catalytic polypeptide	M12679	5.50
Cer1 antigen	NM_014225	5.49
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform	NM_003406	5.49
tyrosine 3-monooxygenase/tyrosinase 5-monooxygenase activator protein, zeta polypeptide	NM_005273	5.45
guanine nucleotide binding protein (G protein), beta polypeptide 2	AF029750	5.43
TAP binding protein (Tapasin)	NM_002394	5.42
soluble carrier family 3 (members of dibasic and neutral amino acid transport), member 2	NM_003145	5.41
signal sequence receptor, beta (translocase-associated protein beta)	U17496	5.36
proteasome (prosome, macropain) subunit, beta type, 8 (large multifunctional protease 7)	NM_016337	5.33
BDG5	NM_000581	5.29
glutathione peroxidase 1	B1570122	5.28
QMA0123 protein	NM_004111	5.27
flag structure-specific endonuclease 1	BC005876	5.27
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 21kD	NM_017528	5.27
patatine methyltransferase	U23946	5.26
RNA binding motif protein 5		

GTP cyclohydrolase I feedback regulatory protein	NM_005258	5.25
12A histone family, member O	A313324	5.25
hypothetical protein B32184_1	BC001648	5.25
larnesyl-diphosphate larnesyltransferase 1	BC003573	5.25
phycocyanin, synaptic 2	NM_004668	5.22
peptide/protein isomerase E (protophilin E)	AF042386	5.21
FKBP6 binding protein 1A (12kD)	BC005147	5.19
guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1	NM_006098	5.15
lymphocyte-specific protein tyrosine kinase	U07236	5.13
MORF melanosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)	NM_006739	5.12
homo box A1	AC004079	5.11
HLA-B associated transcript 3	NM_004639	5.11
tubulin alpha 6	BC005946	5.10
endothelinase 6	NM_004435	5.10
similar to KREB1 cDNA Z31040617 gene	BF972185	5.09
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	M18468	5.09
Consensus includes gbls531931 /FEJAE=gt:13523521 /D0_XREF=est:6025610071 /CLONE=HMAE-4699176 /UG=Hs.108104 ubiquitin-conjugating enzyme E2L 3 /FL=gb	B6531983	5.08
protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	BC000425	5.07
proteasome (prosome, macropain) subunit, beta type, 3	NM_002795	5.07
T cell receptor alpha locus	M12423	5.06
MORF melanosome maintenance deficient 7 (S. cerevisiae)	D55716	5.06
nuclear distribution gene C (A. nidulans) homolog	AF241788	5.05
T cell receptor alpha locus	L34703	5.04
HLA-B associated transcript 3	BC003133	5.02
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), member D	AL566172	5.00
actin, gamma 1	AJ145192	4.99
histon-H2a 4	AC002301	4.98
haryphatin (limpstatin) beta 3	NM_002271	4.98
trehalase of fiver mitochondrial membrane 10 homolog (yeast)	NM_012456	4.98
bacterial 10P repeat-containing 5 (survivin)	A8028869	4.98
stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein)	AL553320	4.97
ribosomal protein, large, P1	NM_001003	4.97
myosin 1D	A4621962	4.97
Zq28.2000bp sequence contig, ORF	B676218	4.96
GTPasease phosphatophylase A	NM_013335	4.96
casein kinase 2, beta polypeptide	NM_001320	4.91
proteasome (prosome, macropain) 35S subunit, non-ATPase, 4	U51007	4.90
cuffin 1	NM_003592	4.90
Hurlerlin interacting protein 8	AF049103	4.90
HLA-E histocompatibility antigen, class I, C	M90684	4.87
solute carrier family 7 (cationic amino acid transporter, y ⁺ system), member 5	A8018009	4.86
meic fringe homolog (Drosophila)	A738965	4.85
RNA-binding protein (leucostigminic)	NM_016732	4.85
heat shock 70MD protein 4	BC002526	4.84
SRF/SRF related, matrix associated, actin dependent regulator of chromatin, subfamily f, member 1	AF231056	4.84
leucine rich repeat (in FLU) interacting protein 1	NM_004735	4.83
Rho GTP dissociation inhibitor (EDI) alpha	BC006337	4.82
adaptor-related protein complex 2, sigma 1 subunit	NM_002341	4.81
lymphocytin beta (TNF superfamily, member 3)	NM_001862	4.81
cytochrome c oxidase subunit Vb	NM_013265	4.79
chromosome 11 open reading frame2	NM_005001	4.78
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5D, 814.5a)	AA502912	4.77
NAA0905 protein	NM_001774	4.77
splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor)	BC004549	4.74
CD37 antigen	B6491442	4.74
tubulin alpha 6	L19164	4.73
block of proliferation 1	AF113682	4.72
peroxiredoxin 1	NM_000075	4.71
thymopressin	BC001675	4.70
cyclic-dependent kinase 4	M15565	4.69
ribosomal protein L13a	BC003049	4.66
T cell receptor alpha locus		
PAI-1 mRNA-binding protein		

DNEZ43401813 protein	B6168666	4.14
c-2 and 5-phase expressed 1	BF305380	4.14
ribosomal protein L27a	NM_000990	4.14
neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	NM_000255	4.13
serum/glucocorticoid regulated kinase-11a	NM_013257	4.13
FGS6 binding protein 1A (120k)	BC001002	4.13
phosphoglucosyl dehydrogenase	NM_002631	4.13
non-POU-domain-containing, octamer-binding	BC003129	4.12
RNA binding motif protein 10	AL137421	4.12
serpinyl antipeptidase	NM_005724	4.12
tetraspin 3	NM_005724	4.12
Enting sarcoma breakpoint region 1	BC006327	4.10
ret finger protein	AF230394	4.10
ribonuclease S precursor	NM_003730	4.09
tumor protein D52	BC389015	4.09
Consensus includes gb:56537190 /FEA=EST /DB_XREF=gi:13528922 /DB_XREF=est:60255589F1 /CLONE=IMAGE-4690079 /UG=Hs.111334 ferritin, light polypeptide	B6537190	4.08
BCL2-interacting killer (apoptosis-inducing)	NM_001197	4.08
RNA310 gene product	BC001404	4.07
accessory proteins BMP31/AMP29	NM_005745	4.06
proteasome (prosome, macropain) 26S subunit, non-ATPase, 9	NM_002813	4.05
neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	AW072388	4.05
butyrophilin, subfamily 3, member A2	BC002832	4.04
B-cell associated protein	NM_007273	4.03
ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 56/58kD, isoform 2	NM_001693	4.02
mitochondrial ribosomal protein S7	NM_015971	4.01
major histocompatibility complex, class I, E	NM_005516	4.00
mitogen-activated protein kinase kinase 3	AA703381	4.00
RNA6599 protein	BC002327	3.99
protein phosphatase 1, regulatory subunit 7	NM_002712	3.99
RNA0729 protein	AW502434	3.98
thyroid autoantigen 70kD (Hu antigen)	NM_001469	3.97
proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)	NM_002800	3.97
hypothetical protein MC5555	NM_024057	3.96
polymyoma (RNA) II (DNA directed) polypeptide 1 (14.5kD)	AL073557	3.94
B-cell translocation gene 1, anti-proliferative	AL535380	3.94
hypothetical protein FL20859	NM_002734	3.93
ATP citrate lyase	U18197	3.92
GAD complex locus	AF064092	3.92
gb:M24663.1 /DEF=Human Ig rearranged H-chain V-region mRNA (c-DJH6), complete cds. /FEA=miRNA /GEN=IGHG /DB_XREF=gi:185200 /FL=gb:M24663.1	M24669	3.92
major histocompatibility complex, class I, E	M31183	3.91
Insulin induced gene 1	NM_005542	3.91
Consensus includes gb:A001784 /FEA=EST /DB_XREF=gi:320255 /DB_XREF=est:041 g06.s1 /CLONE=IMAGE:1619386 /UG=Hs.300332 ESTs. Highly similar to M42735 ribosomal protein M001784	M001784	3.91
Consensus includes gb:282202 /DEF=Human DNA sequence from clone RP1-34P24 on chromosome 22 Contains a pseudogene similar to ribosomal protein L35, E51s, S17s and G5s /FEA=	Z82202	3.89
heterogeneous nuclear ribonucleoprotein R	NM_005826	3.89
platelet activating factor acetylhydrolase, isoform 1b, gamma subunit (29kD)	NM_002573	3.89
hypothetical protein	B6257762	3.88
cytochrome c oxidase subunit Vb	BC066229	3.88
DNEZP547E1010 protein	AF261137	3.88
interferon regulatory factor 3	NM_001571	3.87
postmitotic segregation increased 2-like 9	U38979	3.86
putative cyclin G1 interacting protein	NM_006349	3.86
GMS complex locus	NM_000516	3.86
cyclin-dependent kinase 5	NM_004935	3.86
hepatocarin-like 1	NM_002149	3.86
24-dehydrocholesterol reductase	NM_014762	3.86
2',5'-bisphosphoryl synthetase 1 (40-46 kD)	NM_002534	3.85
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 30	NM_014966	3.85
CREBP/EP200 inhibitory protein 1	AF274951	3.85
adenylate kinase 2	U39945	3.84
structure specific recognition protein 1	BE795648	3.84
Rho GTP dissociation inhibitor (RDI) alpha	BC005851	3.84
Consensus includes gb:J04802 /DEF=Homo sapiens UB8P2 pseudogene for ubiquitin UB8 /FEA=CD5 /DB_XREF=gi:37582 /UG=Hs.247890 Homo sapiens UB8P2 pseudogene for ubiquitin	J04802	3.84
ubiquitin-conjugating enzyme E2H (UBK12 homolog, yeast)	NM_003969	3.83

adenosine deaminase	NM_000022	3.83
basin 2 (60-70kD)	A0284075	3.82
autalytic translation initiation factor 2, subunit 3 (gamma, 52kD)	NM_0016415	3.82
hypothetical protein P101847	AF119855	3.82
enhancer of invasion 10	NM_021178	3.82
HOX2 mitochromosome maintenance deficient 2, mitotin (S. cerevisiae)	NM_004526	3.81
WD repeat domain 1	U63131	3.81
WD repeat domain 1	A8010427	3.81
A kinase (PRK4) anchor protein 2	BE679367	3.81
neural precursor cell expressed, developmentally down-regulated 8	NM_006156	3.80
cathepsin receptor, family 1, subfamily K, member 1	NM_018835	3.80
GMS complex locus	AF088184	3.79
translocase of inner mitochondrial membrane 23 homolog (yeast)	NM_006327	3.79
O-6-methylguanine-DNA methyltransferase	NM_002412	3.79
autalytic translation elongation factor 1 alpha 1	NM_001402	3.79
H3 histone, family 3B (H3.3b)	NM_005324	3.79
RNA974 protein	BE551340	3.78
S-adenosylmethionine decarboxylase 1	M21154	3.77
hypothetical protein M625244	NM_031213	3.76
nucleoside binding protein 2 (RnaD homolog, E. coli)	NM_012225	3.75
cat-delta 1, COAT1 displacement protein (Drosophila)	NM_001913	3.75
splicing factor 3a, subunit 3, 60kD	NM_006802	3.75
ribosomal protein S18	NM_022351	3.74
zinc finger protein 259	NM_003904	3.74
paired immunoglobulin-like receptor beta	NM_013440	3.73
serine/threonine kinase 15	NM_003600	3.73
chromosome 20 open reading frame 1	AF098158	3.73
SH3 domain binding glutamic acid-rich protein like 3	NM_031286	3.72
cyclin G2	L49506	3.72
ANP2 actin-related protein 2 homolog (yeast)	NM_005722	3.71
ubiquitin specific protease 7 (herpes virus-associated)	NM_003470	3.69
deoxyhypusine synthase	NM_001930	3.69
serologically defined colon cancer antigen 16	BC001149	3.69
ribosomal protein L10	AL031276	3.68
PRKX, apoptosis, WT1, regulator	A336206	3.68
hydroxyphenyl-Coenzyme A thiolase(phenyl-Coenzyme A thiolase (bifunctional protein), alpha subunit	U04627	3.67
gla-22-433.1 (REF-H sapiens protein-serine/threonine kinase gene, complete CDS, FEA=mnRNA/P100D=protein-serine/threonine kinase P08_XREF=g:405744 /rL=g:225433.1	Z25433	3.66
RNA-G-related gene	AI167775	3.66
mesenchymal stem cell protein DSC2	NM_016645	3.65
SXT (sets determining region T)-box 2	AF007161	3.65
chromatin-specific transcription elongation factor, 140 kDa subunit	NM_007192	3.65
uncoupling protein 2 (mitochondrial proton carrier)	U62819	3.64
linasin-Gla 5 (mitotic centrosome-associated kinesin)	A1026905	3.63
lymphoid-restricted membrane protein	NM_006152	3.63
T-cell leukemia/lymphoma 1A	BC003574	3.63
ATPase, H+-transporting, lysosomal (vacuolar proton pump) 42kD	NM_0071695	3.63
chromatin assembly factor 1, subunit A (p150)	NM_005483	3.61
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; k)	NM_005789	3.60
IGAM1007 protein	BC000779	3.60
gla maturation factor, gamma	NM_004877	3.59
POP7 (processing of precursor, S. cerevisiae) homolog	BC001430	3.59
breakpoint cluster region protein, uterine leiomyoma, 1; barrier to autointegration factor	A044773	3.59
ubiquitin-Gla 5	NM_024292	3.59
hypothetical protein NF140225	NM_030799	3.58
IBR2	BC000924	3.58
deoxyhypusine synthase	U25266	3.58
peroxiredoxin 2	NM_005809	3.58
Consensus includes gla-8731738 FEA=EST /OB_XREF=g:10145730 /OB_XREF=est:60156815461 /KLONE=IMAGE3842844 /UG=Hs.182937 peptidylprolyl isomerase A (cyclophilin A)	BE731738	3.58
sperm associated antigen 9	NM_003971	3.57
HOX4 mitochromosome maintenance deficient 4 (S. cerevisiae)	A039865	3.57
basin receptor 1 (67kD, ribosomal protein 5A)	AI136306	3.57
vilia 2 (ecoli)	A4670344	3.57
zinc finger protein	NM_015871	3.57

ubiquitin 8	204801	3.56
metastasis-associated 1-like 1	BE222618	3.56
hypothetical protein FU10597	NM_018150	3.54
similar to mouse G83 or D. melanogaster transcription factor IIB	NM_012642	3.54
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; K)	BC026894	3.53
intraflagellar adhesion molecule 3	NM_002162	3.53
hypothetical protein HSC10433	NM_024321	3.53
mitochondrial ribosomal protein S17	NM_015969	3.52
endothelial differentiation-related factor 1	AB022282	3.52
gamma-tubulin complex protein 2	NM_006659	3.51
salivaryferase 1 [beta-galactoside alpha-2,6-sialyltransferase]	AV695711	3.51
transcription factor C72	U03494	3.51
RNA0152 gene product	NM_014730	3.51
Fancet anemia, complementation group A	AF033279	3.51
high-mobility group 208	AZ28679	3.50
zinc finger protein 313	AL031685	3.50
cellular protein elongation factor 1 gamma	AF119850	3.49
nuclear protein family A, member 3 (N/A) small nuclear RNPs	NM_018648	3.48
NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NM_002496	3.48
proteasid protein 2 (colonic epithelium-enriched)	NM_002668	3.48
p47	NM_016143	3.47
RNA0074 protein	D38553	3.47
nuclear distribution gene C (A. nidulans) homolog	NM_006600	3.46
cyclochrome c-1	NM_001916	3.46
Consensus includes gb:574774.1 /REF=g59yn(T) (=OMT3-induced calcium influx regulator human, Jurkat J6 T cell line, mRNA Partial, 1605 nt, /FEA=CD5 /PROD=tyrosine kinase p59yn(T)	574774	3.46
hypothetical protein FU10479	NM_018106	3.46
hypothetical protein FU12287 similar to semaphorins	NM_023367	3.45
apoptosis antagonizing transcription factor	AF083208	3.45
cellular protein elongation factor 1 alpha 1	NM_001403	3.44
ribosomal protein L15	AZ279903	3.44
chromosome condensation 1	NM_001269	3.43
hypothetical protein FU10439	BC006351	3.43
gb:AT014272.1 /REF=Homo sapiens FUS630 mRNA, complete cds, /FEA=mRNA, GEN=FGS630, /PROD=FGS630, /REF=gi:12408251 /UG=Hs.315492 Homo sapiens FUS630, /AV014272	NM_004168	3.43
Consensus includes gb:0027223.1 /REF=Homo sapiens cDNA, FLJ23572 fs, clone UG12403, /FEA=mRNA, /DB_REF=gi:10440306 /UG=Hs.201593 Homo sapiens cDNA, FLJ23572 fs, d	U03743	3.43
SHC [Src homology 2 domain-containing] transforming protein 1	A091079	3.42
succinate dehydrogenase complex, subunit A, flavoprotein (Fp)	NM_004168	3.42
casein kinase 2, beta polypeptide	M30448	3.42
adenylate kinase 2	W02312	3.42
lysine 3-monooxygenase/hydrophobic 5-monooxygenase activation protein, theta polypeptide	NM_006826	3.42
glutamic pyruvate transaminase (alanine aminotransferase)	NM_005309	3.42
histone H4	BF116254	3.42
brucella abortus amplified sequence 4	NM_017843	3.42
ribosomal protein L10	AF057781	3.42
prolactin	NM_002634	3.42
zinc finger protein 183 (RING finger, CHK4 type)	BC000832	3.41
springstone-1 phosphatase	NM_030791	3.41
hypothetical protein MG11256	NM_024324	3.41
nucleophosmin/nucleoplasm 3	NM_006993	3.40
hypothetical protein FU11856	AK021918	3.40
LM and SH3 protein 1	NM_006148	3.40
ribonucleotide reductase M2 polypeptide	BC001886	3.40
casein kinase 1, delta	AV700224	3.39
forkhead box M1	NM_021953	3.39
endoxon 1, (alpha)	NM_001428	3.38
RNA binding protein 7	AL173235	3.38
general transcription factor IIC, polypeptide 3 (102kD)	NM_012086	3.38
postmitotic segregation increased 2-like 11	U38960	3.38
pyridoxal (pyridoxine, vitamin B6) kinase	NM_003681	3.37
hypothetical protein FLJ20340	NM_017773	3.37
alanyl-RNA synthetase	NM_001605	3.37
hurd-like 1 (E. coli)	NM_003707	3.36
Consensus includes gb:8537255 /FEA=EST /DB_REF=gi:13529117 /DB_REF=est:602565318f1 /CLONE=IMAGE:4689748 /UG=Hs.82646 Dnal (Hsp40) Homolog, subfamily B, member	8537255	3.36

Ras-GRTPase activating protein SH3 domain-binding protein 2	NM_012297	3.36
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1	AK021419	3.35
hypothetical protein [MAGE3455200]	NM_024006	3.34
cupping protein [actin filament] muscle 2-like, beta	NM_004930	3.34
Dual (Hsp40) homolog, subfamily C, member 8	AA643895	3.34
transportin-SR	NM_012470	3.34
oxidase (cytochrome c) assembly 1-like	BC001663	3.33
Sigma3 synthase nuclear activator 1	BC004118	3.33
protein synthetase co-transcribed (bacterial homolog)	NM_007198	3.33
interleukin stimulator gene (IL20)	U88564	3.32
protease (prosome, macropain) subunit, beta type, 4	NM_002796	3.32
serine hydroxymethyltransferase 2 (mitochondrial)	AB750316	3.32
HPF2 non-histone chromosome protein 2-class 1 (S. cerevisiae)	NM_005008	3.32
serine/threonine kinase 12	AB011446	3.32
COP9 homolog	NM_006710	3.32
HSPX160 protein	NM_014182	3.31
nuclear matrix protein HMY200 related to splicing factor PRP19	NM_014502	3.31
myosin, light polypeptide 6, alpha, smooth muscle and non-muscle	BE724356	3.31
ATX1 antioxidant protein 1 homolog (yeast)	NM_004045	3.31
MTX1 protein	NM_021640	3.31
hydrophobic membrane protein	U10485	3.30
PD RNA-binding protein	L03411	3.29
histidyl-tRNA synthetase	NM_002109	3.29
hypothetical protein MG2650	NM_024108	3.29
adenine phosphoribosyltransferase	AA027724	3.28
BDJ23 homolog A (S. cerevisiae)	BF572938	3.28
g04274954.1 (DEF=Homo sapiens PNAS-29 mRNA, complete cds. /FEA=mRNA /PID=PNAS-29 /IDL_XREF=gi:12751070 /UG=Hs.326791 Homo sapiens PNAS-29 mRNA, complete cds. /)	NM_024954	3.28
hypothetical protein FUJ2549	NM_024844	3.28
immunoglobulin lambda locus	AF047245	3.27
hypothetical protein FUJ1305	NM_018386	3.27
HSPC002 protein	NM_015362	3.27
transin	AF35302	3.25
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	M65254	3.24
death associated protein 3	U18321	3.24
SHG-interacting protein	AF689564	3.24
hypothetical protein FLJ20552	NM_017876	3.23
uncharacterized hypothalamus protein HT009	NM_018470	3.23
mitochondrial ribosomal protein L12	NM_002949	3.23
memosides, alpha, class 1A, member 1	NM_005907	3.23
histone H2A17Z variant	BF343852	3.23
dihydropyrimidine 5-acyltransferase (E2 component of pyruvate dehydrogenase complex)	J03866	3.23
ATP-dependent interferon responsive	NM_022371	3.23
nucl5 homolog 5 (E. coli)	AF034759	3.23
X-prolyl aminopeptidase (aminopeptidase P)-like	NM_006523	3.22
villin 2 (catn)	J05021	3.22
map kinase phosphatase-like protein MK-STYX	NM_016086	3.22
receptor II	BC003560	3.22
KAM1608 protein	NM_024820	3.21
G3 protein	NM_016947	3.21
eukaryotic translation initiation factor 2B, subunit 2 (beta, 39kD)	NM_014239	3.21
nuclear transcription factor Y, gamma	U62296	3.20
Integral type I protein	BC000027	3.20
ATP synthase, H ⁺ -transporting, mitochondrial F1 complex, beta polypeptide	NM_001686	3.20
Consensus includes g04025759.1 (DEF=Homo sapiens cDNA: FLJ22106 fs, clone HEP17675. /FEA=mRNA /IDL_XREF=gi:10438374 /UG=Hs.323470 Homo sapiens cDNA: FLJ22106 fs, d	AK025759	3.20
dynamin, cytoplasmic, heavy polypeptide 1	AB002323	3.20
change and polyadenylation specific factor 1, 16kD subunit	NM_013291	3.20
nucleic [nucleoside diphosphate linked moiety X]-type motif 1	NM_002452	3.19
hypothetical protein FUJ0374	NM_018074	3.19
glutamic-catalase-like transferase 2, mitochondrial (separate aminotransferase 2)	NM_002080	3.18
candidate tumor suppressor protein	NM_020381	3.18
KAM1049 protein	AF322111	3.17
Consensus includes g04118510 (DEF=Human DNA sequence from clone RPS-858M22 on chromosome 20. Contains a pseudogene similar to 40S ribosomal protein S10, STSs and G55a /FEA=AL118510	AF322111	3.17
tumor suppressor deleted in oral cancer-related 1	NM_005851	3.16

phosphoserine isomerase 1	NH_000385	3.16
phosphoribosyl pyrophosphate synthetase 1	NH_002764	3.16
aldo-keto reductase family 1, member A1 (aldehyde reductase)	NH_006066	3.15
major histocompatibility complex, class II, DR beta 5	NH_002125	3.15
aryl hydrocarbon receptor interacting protein	AL58532	3.15
prostate tumor over expressed gene 1	AL046054	3.15
anaphase promoting complex subunit 5	BF967271	3.15
pepck1, peck1 isomerase H (pyruvate H)	NH_006347	3.14
major histocompatibility complex, class I, C	AK024836	3.14
adenylosuccinate lyase	NH_000026	3.13
adaptor protein	U78525	3.13
adaptor protein	NH_020187	3.13
cell binding protein	NH_014593	3.13
divalent cation tolerant protein (CUTA)	AF230924	3.12
hypothetical protein	AA243508	3.12
NAOH dehydrogenase (ubiquinone) Fe-S protein 6 (13UD) (NAOH-coenzyme Q reductase)	NH_004553	3.12
Consensus includes gb:8547674 /FEA=EST /DB_XREF=gi:9776319 /DB_XREF=est:60107625551 /CLONE=IMAGE:3462325 /JG=Hs_204169 ESTs	BE547674	3.11
hypothetical protein similar to mouse aldehyde reductase 6 (renal)	AL096767	3.11
GMP22 for guanine monophosphate reductase isolog	NH_016576	3.11
vesicle-associated membrane protein 1 (GlypDomein 1)	NH_016530	3.11
acetonamide nucleotide transferase	NH_012343	3.10
cell division cycle 2, G1 to S and G2 to M	NH_007786	3.10
hypothetical protein FLJ12660	NH_025152	3.10
BAG7, member RAS oncogene family	AK000826	3.10
nucleoside phosphorylase	NH_000270	3.10
sorting nexin 5	NH_014426	3.09
core histone macroH2A.2	NH_018649	3.09
adaptor protein	AB044548	3.08
adaptor protein	NH_001419	3.08
ELAV (elav-type leish, abnormal elav, Oncoapla)-like 1 (Hu antigen B)	BC053773	3.08
mitochondrial ribosomal protein S18B	NH_014303	3.07
peccadillo homolog 1, containing BKT domain (zebrafish)	NH_001494	3.07
GTP dissociation inhibitor 2	U28936	3.07
lysine 3-monoxygenase/hydrophase 5-monoxygenase activation protein, epsilon polypeptide	NH_018031	3.07
BD repeat domain 6	NH_004719	3.07
splicing factor, arginine/serine-rich 2, interacting protein	AK022732	3.07
hypothetical protein FLJ20436	NH_004130	3.07
glycoprotein	AB03180	3.07
phosphorine aminotransferase	NH_001363	3.07
dyskeratosis congenita 1, dyskerin	NH_001281	3.07
cytoskeleton-associated protein 1	AI91427	3.07
neural precursor cell expressed, developmentally down-regulated 5	NH_000386	3.06
bleomycin hydrolase	NH_004939	3.06
DEAD/H (Asp-Glu-Ala-Gly/His) box polypeptide 1	NH_003105	3.06
serpin-related receptor, L (DLR class) A repeats-containing	L40326	3.06
damage-specific DNA binding protein 1 (127AD)	Z97353	3.06
ribosomal protein L15	NH_022156	3.05
PP3111 protein	AF05484	3.05
sorting nexin 1	AK027546	3.05
zinc finger protein 143 (clone pHz-1)	NH_001328	3.05
C-terminal binding protein 1	NH_018292	3.05
hypothetical protein FLJ10989	B25889	3.05
ULK domain-containing 2	AB030314	3.04
ribosomal protein S2	S67779	3.04
B-cell CLL/lymphoma 6 (zinc finger protein 51)	NH_004317	3.04
avr4 (bacterial) serine transporters, ATP-binding, homolog 1	NH_005496	3.04
SHC4 structural maintenance of chromosomes 4-like 1 (Yeast)	BE568833	3.04
Consensus includes gb:8564833 /FEA=EST /DB_XREF=gi:10575538 /DB_XREF=IMAGE:3933782 /JG=Hs_324648 Homo sapiens cDNA FLJ13700 fts, clone PLA	NH_018128	3.04
hypothetical protein FLJ10534	NH_004470	3.04
PTC06 binding protein 2 (13AD)	NH_001629	3.03
archoninase 5-lysozyme-activating protein	AF141970	3.03
interleukin enhancer binding factor 3, 300D	NH_001175	3.03
the GTP dissociation inhibitor (GDI) beta	D86973	3.03
GDI1 general control of amino-acid synthesis 1-like 1 (Yeast)		

replication factor C (activator 1) 5 (S6.540)	NM_007370	2.89
polymerase (DNA-directed), delta 4	NM_021173	2.89
arc family associated phosphoprotein 2	AC03999	2.88
presenilin 1 (Alzheimer disease 3)	NM_007318	2.88
cytoskeleton-associated protein 1	BC005969	2.88
integrin, alpha 4 (antigen CD490, alpha 4 subunit of VLA-4 receptor)	NM_000885	2.88
NAA1228 protein	A708767	2.88
mannosidase, alpha class 2B, member 1	U68567	2.88
ADP-ribosylation factor 4	AL537042	2.87
GLA-150 protein	AF061730	2.87
hypothetical protein M6ZG555	NM_024339	2.87
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	NM_002716	2.87
interleukin 21 receptor	NM_021798	2.87
thyroid receptor interacting protein 15	NM_004236	2.87
splicing factor, arginine/serine-rich 7 (3500)	AC524053	2.87
ras homolog gene family, member H	NM_004310	2.87
SEC13-like 1 (S. cerevisiae)	NM_030673	2.86
CD4-MOT transcription complex, subunit 3	NM_014516	2.86
sporadic sarcoma translocation, chromosome 18	NM_005637	2.86
NAA1513 protein	NM_017542	2.86
FGS06 binding protein 1A (F202)	A936769	2.86
PRF2 periodic tryptophan protein homolog (yeast)	U56085	2.86
HSP134 protein	NM_014169	2.86
transcriptional adaptor 3 (ADA3, yeast homolog-like (PCAF histone acetylase complex)	A024982	2.85
transferrin receptor (990, CD71)	NM_003234	2.85
ubiquitin factor E48 (UFD2 homolog, yeast)	A028839	2.85
Williams-Beuren syndrome chromosome region 5	A7257135	2.85
branched chain aminotransferase 1, cytosolic	A652662	2.84
hypothetical protein	NM_015684	2.84
delonin A, fructose-bisphosphate	A026577	2.84
maculoid-like (Drosophila)	N31913	2.84
GLA-120 protein	NM_016057	2.83
selective kinase	U95339	2.83
HLA-E histocompatibility antigen, class I, 6	M96886	2.83
Consensus includes gp:AM68D502 PEA=EST /DB_XREF=gi:2556270 DB_XREF=est:ac83405.s.1 /CLONE=IMAGE589193 /UG=Hs.181125 immunoglobulin lambda locus	A680302	2.83
high-mobility group (nonhistone chromosomal) protein 14	NM_004965	2.83
gdy3 associated, gamma adaptin ear containing, ANF binding protein 1	NM_013365	2.83
Consensus includes gp:AL353641 DEF=Humer DNA sequence from clone RP1-158P9 on chromosome 1. Contains a putative novel gene, a laminin receptor 1 (6740, ribosomal protein 5A) AL353641	NM_012392	2.82
PEF protein with a long N-terminal hydrophobic domain (peffin)	NM_016050	2.82
mitochondrial ribosomal protein L11	BC001149	2.81
serologically defined colon cancer antigen 16	NM_001985	2.81
electron-transfer flavoprotein, beta polypeptide	X06130	2.81
chromosome condensation 1	AF083389	2.80
Ref-Hirschhorn syndrome candidate 1	AF131863	2.80
SKD-domain, 6182-84a, endoplasmic R	A044944	2.80
mitochondrial ribosomal protein S11	BC001416	2.80
protein phosphatase 4 (formerly X), catalytic subunit	NM_016613	2.79
AD021 protein	AA974416	2.79
protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform	NM_007023	2.79
similar to S. cerevisiae RER1	BC474429	2.79
putative homolog 1 (Drosophila)	AF591354	2.79
heterogeneous nuclear ribonucleoprotein F	AF04206	2.79
inhibitor of growth family, member Y	AF040076	2.78
for (Pro/His related) tyrosine kinase (phosphoprotein MCF94)	NM_005246	2.78
Francis aments, complementation group 6	AL004472	2.78
nuclear receptor subfamily 3, group C, member 1	U01351	2.78
immunoglobulin lambda locus	X33066	2.78
ATPase, H+-transporting, lysosomal (vacuolar proton pump) 16A0	NM_001694	2.78
gdy3 transport complex 1 (90 kDa subunit)	NM_006348	2.78
hypothetical protein FLJ22940	NM_024571	2.77
apoptosis related protein APR-3	NM_016085	2.77
RNA0095 gene product	NM_014669	2.77

synaptophysin-like protein
 G protein-coupled receptor 18
 hypothetical protein PF01635
 gpAF008937.1 DEF=Homo sapiens syntaxin-16c mRNA, complete cds. /FEA=mRNA PRODB=Homo sapiens syntaxin-16c /DB_XREF=gi:2352817 /JGI=Hs.102178 syntxin 16 /FL=gpAF008937.1
 hypothetical protein FL20517
 ribosomal protein L5
 proteinase (prosome, macroquin) subunit, beta type, 10
 hypothetical protein NC22722
 chromosome 11 open reading frame 9
 hypothetical protein FLJ12442
 CD164 antigen, sialomucin
 actinin 1
 glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1)
 arginine-glutamic acid dehydrogenase (RGE) repeats
 hypothetical protein FLJ14040
 chromosome 11 open reading frame 10
 PPM4/STK10 splicing factor
 transformation/transcription domain-associated protein
 RIMM999 protein
 self-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phosphoperoxidase A2)
 reticoid control protein ds3 homolog
 DNA segment, single copy probe LUG-OU/UG-CAT (deleted in polyposis)
 boottitis dehydrogenase 3 (NUO-3) beta
 SMY3 suppressor of mit two 3 homolog 1 (yeast)
 translocase of outer mitochondrial membrane 20 (yeast) homolog
 ribosomal protein L24
 UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylglucosaminyltransferase 2 (GALNAc-T2)
 BT33 associated nuclear protein
 hypothetical protein FLJ12298
 ribosomal protein S10
 RAGA, member RAS oncogene family
 RAGM731 protein
 boottitis dehydrogenase 3 (NUO-3) beta
 RAGM144 gene product
 v-crk avian sarcoma virus CT10 oncogene homolog
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1
 inhibitor of DNA binding 3, dominant negative helix-loop-helix protein
 hypothetical protein FLJ22837
 methionine-tRNA synthetase
 hypothetical protein FLJ22795
 DEAD/H (Arg-Glu-Ala-Asp/Glu) box polypeptide 35
 glutathione peroxidase 4 (phospholipid hydroperoxidase)
 hypothetical protein FLJ11301
 Consensus includes gp:BP796470 /FEA=EST /DB_XREF=gi:12101524 /DB_XREF=est:60225992.6F1 /FLONE=Hs.4342999 /JGI=Hs.284275 Homo sapiens PAN2 mRNA, complete cds. /FL=FL136897
 5-phosphoglucomutase
 poly(A) binding protein, nuclear 1
 polymerase (RNA) II (DNA directed) polypeptide D
 proliferation-associated 254, 3880
 RAGM542 gene product
 thymidyl-RNA synthetase
 RDEL (Lys-Arg-Glu-Leu) endoplasmic reticulum protein retention receptor 1
 phosphatase 1
 putative protein similar to heavy (tropomyosin)
 cytochrome P450, subfamily 51 pseudogene 2
 c-est tyrosine kinase
 ATPase, Na+/K+ transporting, beta 3 polypeptide
 BUB3 binding uninhibited by benzimidazoles 3 homolog (yeast)
 peroxiredoxin 2
 quinoid dihydropteridine reductase
 mitochondrial carrier homolog 1
 protein kinase C and casein kinase substrate in neurons 2
 hypothetical protein FLJ11301
 putative nuclear protein

hypothetical protein NM_013298 2.68
chromodomain protein, Y chromosome-like AF061258 2.68
Cluster IncL AI131332x0800.1 Homo sapiens cDNA, 3' end /clone=IMAGE:567097 /clone_end=3 /gi=AA131335 /gi=1692833 /ug=Hs.28166 /len=552
carboxylate (N-acetylglucosamine 6-O) sulfotransferase 6 AA131335 2.68
Consensus includes gb:BE878277 /FEA=EST /DB_XREF=gi:10327053 /DB_XREF=est:601488188F1 /CLONE=IMAGE:3890564 /UC=Hs.57079 Homo sapiens cDNA FL13267 fs, clone OVA1 BE878277 NM_021615 2.67
phosphatidylinositol glycan, class F NM_021615 2.67
ubiquitin-conjugating enzyme E2 variant 1 NM_002643 2.67
hypothetical protein FLJ10774 U93361 2.67
SCAN domain-containing 1 U93361 2.67
CD38 antigen [p45] NM_015904 2.66
microtubule-associated protein 1A/18 light chain 3 NM_015904 2.66
translation initiation factor IF2 NM_015904 2.66
DNA segment on chromosome X (unique) 9879 expressed sequence NM_006014 2.66
mink enzyme 2, MO(+)-dependent, mitochondrial M53505 2.66
ubiquitin-conjugating enzyme E2N (Ubc13 homolog, yeast) B262760 2.66
protease, serine, 15 U02389 2.66
major histocompatibility complex, class I, A U02389 2.66
synectin 7 AA573862 2.65
interferon-related developmental regulator 1 N32035 2.65
ICF-2 mRNA-binding protein 3 NM_001550 2.65
AIP7 actin-related protein 3 homolog (yeast) AU160004 2.65
hepatocellular carcinoma-associated antigen 66 NM_005721 2.65
peptidylprolyl isomerase A (cyclophilin A) NM_018428 2.65
hypothetical protein BC005982 2.64
thrombin domain-containing NM_018986 2.64
major histocompatibility complex, class I, C NM_030755 2.64
TNF-induced protein BC004489 2.64
amyloid beta (A4) precursor-like protein 2 BC005352 2.64
adaptor-related protein complex 2, alpha 2 subunit B4427393 2.64
fibroblast growth factor (bFGF) intracellular binding protein A125280 2.64
voltage-dependent anion channel 3 NM_004214 2.63
ribosomal protein S10 U99943 2.63
chromatin-10 factor 1 NM_001014 2.63
Consensus includes gb:BE93164 /FEA=EST /DB_XREF=gi:1765582 /DB_XREF=est:601656973B1 /CLONE=IMAGE:3865650 /UC=Hs.2186 eukaryotic translation elongation factor 1 gamma BE93164 2.63
postmeiotic segregation increased 2-like 3 NM_006384 2.62
calcium and integrin binding protein (DM-dependent protein kinase interacting protein) NM_004127 2.62
G protein pathway suppressor 1 BC002594 2.62
diallyl-/alpha-phosphatidylglycerol-4-phosphate-protein glycoyltransferase NM_014166 2.62
HSPK126 protein AF116273 2.62
BCL2-associated athanogene BC006301 2.62
anaphase promoting complex subunit 5 NM_006197 2.62
pericentriolar material 1 AF258341 2.61
P450 (cytochrome) oxidoreductase NM_003683 2.61
DNA segment on chromosome 21 (unique) 2056 expressed sequence NM_004030 2.61
interferon regulatory factor 7 AF050641 2.61
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9 (39kD) U93398 2.61
G protein-coupled receptor 68 NM_006527 2.60
stem-loop (histone) binding protein NM_013336 2.60
protein transport protein SEC61 alpha subunit isoform 1 NM_001749 2.60
caspin, small subunit 1 X63465 2.60
MAP1, GTP-ADP dissociation stimulator 1 575762 2.60
fusion, derived from t(12;16) malignant liposarcoma A1962933 2.60
heat shock 90kD protein 1, alpha NM_017601 2.60
hypothetical protein DNG2p761H221 U79718 2.60
nif endonuclease II-like 1 (E. coli) NM_006893 2.59
Segin NM_005174 2.59
ATP synthase, H⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1 NM_005174 2.59
Consensus includes gb:BE544663 /FEA=EST /DB_XREF=gi:3773308 /DB_XREF=est:601074131F1 /CLONE=IMAGE:3460398 /UC=Hs.110407 ESTs, Weakly similar to coded for by C. elegans BE544663 2.59
mesenchymal stem cell protein USK43 A1992095 2.59
BCL2-associated athanogene NM_004323 2.59
transcription factor Dp-2 (E2F dimerization partner 2) BC034328 2.59
gb:BC005668.1 /DBF=Homo sapiens. Similar to zinc finger protein 304, clone MGC:4079, mRNA, complete cds. /FEA=miRNA PR000=Similar to zinc finger protein 304 /DB_XREF=gi:135434 BC005668 2.59
nuclear and coiled-body phosphoprotein 1 NM_004741 2.59

soluble carrier family 25 (mitochondrial carrier, citrate transporter), member 1	U25147	2.58
hypothetical protein FU10769	NM_018210	2.58
TIA1 cytosolic granule-associated RNA binding protein	BF692742	2.58
GMP responsive element modulator	NM_001881	2.58
ubiquitin specific protease 23	A4621731	2.58
hypothetical protein FU10702	NM_018184	2.58
UDP glycosyltransferase 2 family, polypeptide B15	NM_001076	2.57
polymerase (RNA II (DNA directed)) polypeptide 6	NM_002896	2.57
small glutamine-rich tetrapeptide repeat (TPR)-containing	NM_003021	2.57
CG-56 protein	AL022316	2.57
transmembrane 7 superfamily member 2	AF086304	2.57
DISE2756-10123 protein	BC002873	2.57
peroxylhydroxyl isomerase A (cydophilin A)	NM_021130	2.57
RNA0625 protein	N64643	2.57
jumping translocation breakpoint	BC004239	2.56
sub-associated polypeptide, 30kD	BF247098	2.56
cell division cycle 2, G1 to S and G2 to M	088357	2.56
RNA06906 protein	AA502912	2.56
RA021 homolog (S. pombe)	BC289967	2.56
branched chain aminotransferase 2, mitochondrial	NM_001190	2.56
peroxyl-his homodomain transcription factor 1	NM_002653	2.56
hematopoietic cell-specific Lyn substrate 1	NM_005335	2.56
membrane component, chromosome 11, surface marker 1	NM_005898	2.56
anesthetics resistance protein ARS2	NM_015908	2.56
cleavage and polyadenylation specific factor 1, 160kD subunit	U37012	2.56
hypothetical protein FU02044	BC002492	2.55
G1 to S phase transition 1	NM_002094	2.55
RNA0233 gene product	NM_014745	2.55
similar to S. pombe dim1+	NM_006701	2.55
MAO1 mitotic arrest deficient-like 1 (yeast)	NM_003550	2.55
adrenic phosphotransferase	NM_000485	2.55
Consensus includes g0130250, FFA=EST, /B8_38EF=gi:11255432, /B8_38EF=est:6018903081, /CLONE=IMAGE:4131673, /UG=Hs.315103, ESTs. Moderately similar to DNA-binding pro	BF308250	2.55
arginase-rich, related in early stage tumors	NM_006610	2.55
antifolate 21 receptor	AF269133	2.55
clathrin, heavy polypeptide (Hc)	NM_004859	2.55
ubiquitin specific protease 24	AB028980	2.54
immunoglobulin lambda locus	AF043566	2.54
programmed cell death 8 (apoptosis-inducing factor)	NM_004208	2.54
RAE1 RNA export 1 homolog (S. pombe)	U85943	2.54
isolectin (lyso)-1 (or 4)-mannophosphatase 2	NM_014214	2.54
protein regulator of cytokinesis 1	NM_003581	2.54
hypothetical protein FU12973	NM_024908	2.54
RNA0088 protein	W87689	2.54
NSPC023 protein	NM_014047	2.54
vitin 2 (ecoli)	AF199015	2.54
PAI-1 mRNA-binding protein	NM_015640	2.54
RNA0635 gene product	AB022669	2.53
pseudouridylyl synthase 1	NM_025215	2.53
RNA0819 protein	AB020626	2.53
chromosome 12 open reading frame 8	NM_006817	2.53
stand regulatory element binding transcription factor 2	BE513151	2.52
Wol-Hirschhorn syndrome candidate 1	BF111870	2.52
casin lipase 2, alpha prime polypeptide	NM_001896	2.52
Consensus includes g014708130, FFA=EST, /B8_38EF=gi:10725395, /B8_38EF=est:4514708130, /CLONE=LOC40405, /UG=Hs.283965 Homo sapiens scavenger receptor class B type III SR-I	AF043290	2.52
Fea1 homolog 2 (E. coli)	NM_014164	2.52
synaptic nucle expressed gene 1b	NM_018358	2.52
FXR1 domain-containing ion transport regulator 5	AB009358	2.51
hypothetical protein FU11198	NM_002817	2.51
mitogen-activated protein kinase kinase 7	BF663141	2.51
proteasome (prosome, macropain) 26S subunit, non-ATPase, 13	AF758755	2.51
vitin 2 (ecoli)	NM_004781	2.50
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma, Ki)		
vesicle-associated membrane protein 3 (clathrovin)		

chromosome 2 open reading frame 3	AC005034	2.50
ribosomal protein S25	AA883388	2.50
arsenite resistance protein ARS2	BE546076	2.50
a disintegrin and metalloproteinase domain 10	AU135154	2.50
RNA11228 protein	EC001224	2.50
intran light chain 1 slow a	NM_002475	2.50
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g	AL050277	2.50
mitochondrial ribosomal protein L22	NM_014180	2.49
heterogeneous nuclear ribonucleoprotein H3 (H3)	AF132363	2.49
starlin, RNA binding protein (Drosophila)	AI136601	2.49
polypyridine tract binding protein (heterogeneous nuclear ribonucleoprotein 1)	X52006	2.49
S-phase kinase-associated protein 2 (p45)	BC001441	2.49
estrogen-related receptor alpha	L36487	2.49
FKBP5 binding protein 1A (1260)	NM_000801	2.49
hypothetical protein DMT2p56A2022	NM_030796	2.49
regulator of inosine transcripts 1	066988	2.49
HSP162 protein	NM_014183	2.49
CXK6 cell division cycle 6 homolog (S. cerevisiae)	NM_001254	2.49
gRNO_004602.1 / DUF=Homo sapiens PK4 and SFPS1 interacting protein 2 / DUF_XREF=g-4758869	NM_004602	2.48
glyoxylate reductase/hydroxypyruvate reductase	AG026752	2.48
chaperonin containing TCP1, subunit 8 (beta)	NM_006595	2.48
hypothetical gene H6C19595	W67887	2.48
aldo-keto reductase family 1, member 81 (aldose reductase)	NM_001628	2.48
tyrosine 3-monooxygenase/tyrosine 5-monooxygenase activation protein, zeta polypeptide	U28964	2.48
Barak-like growth factor receptor 1 (fms-related tyrosine kinase 2, Phifer syndrome)	NM_023107	2.48
low density lipoprotein receptor (familial hypercholesterolemia)	NM_000527	2.47
HSPC015 protein	NM_014026	2.47
MDZ3 homolog 8 (S. cerevisiae)	AL527365	2.47
RNA binding motif protein 14	NM_006328	2.47
C64-96 protein	AL023316	2.47
arrestin A2	NM_004039	2.47
lamin B2	M94363	2.47
hypothetical protein NCIA-iso	NM_030937	2.47
lactate dehydrogenase C3P1	NM_021633	2.46
RNA11856 protein	AN026933	2.46
MAP kinase-kinase-kinase serine/threonine kinase 1	BC022755	2.46
chromium-activated protein kinase kinase kinase 7	AF218074	2.46
HLA-E histocompatibility antigen, class I, E	M98685	2.46
secretory carrier membrane protein 3	NM_005698	2.46
galactosidase, beta 1	NM_004404	2.46
homolog of yeast Hase HRP/AtHase P protein Pop5	NM_015918	2.45
keratopharin (keratin) beta 1	AC004941	2.45
immunoglobulin lambda3 joining 3	X57812	2.45
gRNM_006326.1 / DUF=Homo sapiens seven transmembrane domain protein (NITE14), mRNA / FEA=mRNA / EN=NITE14 / PROD=seven transmembrane domain protein / DUF_XREF=g-54	NM_006326	2.45
proteasome (prosome, macropain) 26S subunit, ATPase, 1	NM_002802	2.45
hypothetical protein DMT2p43422216	AC004475	2.45
ribosomal protein S15a	NM_001019	2.44
general transcription factor II, 1	NM_001518	2.44
ubiquitin-conjugating enzyme E2D 3 (UBC475 homolog, yeast)	NM_003340	2.44
ribonuclease H1	BC534527	2.44
Consensus includes gR8332462 / FEA=EST / DUF_XREF=g-13138900 / DUF_XREF=est-602432371 / K1ONE=IMAGE4550505 / UG=Hs.3343 phosphoglycerate dehydrogenase	BC532462	2.44
lipic acid synthetase	BF056778	2.44
general transcription factor IIR, polypeptide 4 (52KD subunit)	NM_001517	2.44
hypothetical protein FLU0360	NM_017782	2.44
RNA11028 protein	AB028551	2.44
MDH1 dehydrogenase (ubiquinone) 1 alpha subcomplex, 4 (90D, MLRQ)	NM_002489	2.43
ribosomal protein L7a	NM_000972	2.43
threonine and one amino acid protein kinase	AK007163	2.43
Brk finger protein Z25	AC074331	2.43
TM6 family member-associated H708 activator	NM_004180	2.43
carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydrogenase	NM_004341	2.43
Tad-Hirschhorn syndrome candidate 1	BE793789	2.43
MDL18 protein	AW248552	2.43

nucleotide binding protein
cell division cycle 2-like 2
CDL18 protein
tumor necrosis factor receptor superfamily, member 7
3'-phosphoadenosine 5'-phosphatidate synthase 1
guanine nucleotide binding protein (G protein), alpha 13
angptl-1 interacting protein 1; actin receptor interacting protein 1
hypothetical protein FLJ12806
CCR4-NOT transcription complex, subunit 8
Consensus includes gp:AL365505 /REF=human DNA sequence from clone RP11-382A12 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 5' part of the gene for AL365505
gp:J13287.1 /REF=human mRNA for alternative splicing product of glucocorticoidase pseudogene, complete cds. /FEA=mRNA /PRD=glucocorticoidase /DB_XREF=gi:496370 /UG=Hs.013287
cristin kinase 1, delta
serine/threonine kinase 24 (STE20 homolog, yeast)
pyridine-5-carboxylate synthetase (glutamate gamma-semialdehyde synthetase)
Consensus includes gp:AA485687 /FEA=EST /DB_XREF=gi:2216118 /DB_XREF=est:ab38033.s1 /CLONE=IMAGE:643101 /UG=Hs.284235 ESTs
leucine-rich PPR-motif containing
phosphatidylcholine, plasmalogen
DnaI (Hsp46) homolog, subfamily B, member 6
heterogeneous nuclear ribonucleoprotein C (C1/C2)
desmin (actin depolymerizing factor)
WW domain binding protein 11
gp:NM_016402.1 /REF=human sapiens hypothetical protein (HSPC140), mRNA. /FEA=mRNA /GEN=HSPC140 /PRD=HSPC140 /DB_XREF=gi:7705472 /UG=Hs.279868 hypothetical
HSPC053 protein
malate dehydrogenase 1, MD (soluble)
H2A histone family, member Y
nucleolar and coiled-body phosphoprotein 1
Mei56 (D, melanocortin) like protein
gp:NM_014128.1 /REF=human sapiens PRD0470 protein (PRD0470), mRNA. /FEA=mRNA /GEN=PRD0470 /PRD=PRD0470 protein /DB_XREF=gi:7662555 /UG=Hs.278937 PRD0470 prot
hypothetical protein FLJ10083
DUF25660120 protein
hypopharynx-1 RNA synthetase
adaptor-related protein complex 2, beta 1 subunit
general transcription factor IIF, polypeptide 1 (TAD subunit)
SEC14-like 1 (S. cerevisiae)
cytochrome b5 outer mitochondrial membrane precursor
D-dephospho luciferase
gp:NM_020342.1 /REF=human sapiens PRSET domain containing protein 07 (SET07), mRNA. /FEA=mRNA /GEN=SET07 /PRD=PRSET domain containing protein 07 /DB_XREF=gi:99664854 /NM_020342
synthase 10
Consensus includes gp:H97931 /FEA=EST /DB_XREF=gi:1118816 /DB_XREF=est:tp04c02.s1 /CLONE=IMAGE:260738 /UG=Hs.173108 Homo sapiens cDNA: FLJ1897 fs, clone HEP03447
protein kinase, cAMP-dependent, catalytic, beta
peroxisomal membrane protein 2 (ZEB2)
interleukin 10 receptor, beta
hypothetical protein FLJ10603
Consensus includes gp:86249599 /FEA=EST /DB_XREF=gi:12759426 /DB_XREF=est:605319652F1 /CLONE=IMAGE:4414895 /UG=Hs.110771 Homo sapiens cDNA: FLJ1904 fs, clone HE
antigen identified by monoclonal antibody 16-67
GyrB-like protein coxapene
hypothetical protein FLJ20257
JAK binding protein
hypothetical protein FLJ11773
mitotic spindle coiled-coil related protein
CDC45 cell division cycle 45-like (S. cerevisiae)
interferon-related developmental regulator 2
NOL18 protein
nucleoside SMO
tyrosyl-DNA synthetase
diphthamox homolog 1 (Drosophila)
thymosin, beta 10
tubulin p50 (TUBA0-containing) subunit B 1
cathepsin light polypeptide (Lc4)
cell division cycle 2-like 2
hypothetical protein FLJ21617
cat eye syndrome chromosome region, candidate 5
Consensus includes gp:AB67738 /FEA=EST /DB_XREF=gi:4899032 /DB_XREF=est:tp3g08.s1 /CLONE=IMAGE:2206910 /UG=Hs.659 pepitidylprolyl isomerase B (cyclophilin B)

Consensus includes gb:AL137162 /DEF=Human DNA sequence from clone RPS-843L14 on chromosome 20. Contains ESTs, STSs and GSSs. Contains a novel gene and the 5' part of a gene to AL137162	2.29
gM060511.1 /DEF=Homo sapiens clone 016s10 My016 protein mRNA, complete cds. /FEA=mRNA /PROID=My016 protein /DB_XREF=gi:12001971 /UG=Hs.161634 Homo sapiens cDNA; NM_014630	2.29
RNAQ211 gene product	2.29
isochitrate dehydrogenase 3 (NAD+) alpha	2.29
MD-1, BP105-associated	2.29
transforming growth factor, beta 2	2.29
ring finger protein 10	2.29
gH4N_016007.1 /DEF=Homo sapiens (LOC51625), mRNA. /FEA=mRNA. KEN=LOC51625 /PROID=C61-S9 protein /DB_XREF=gi:7706297 /UG=Hs.279867 CG1-S9 protein /FI NM_016007	2.29
ORF274340156 protein	2.28
hypothetical protein NC02731	2.28
acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase)	2.28
glutaryl-prolyl-4RNA synthetase	2.28
Consensus includes gb:AK000185.1 /DEF=Homo sapiens cDNA FLJ20178 fs, clone COL09990. /FEA=mRNA /DB_XREF=gi:7020104 /UG=Hs.306389 Homo sapiens cDNA FLJ20178 fs, clone AK000185	2.28
hypothetical protein FLJ20764	2.28
protease, serine, 25	2.28
mitogen-activated protein kinase kinase kinase 12	2.28
RSL1-Ela (Yeast)	2.28
regulator of G-protein signalling 10	2.28
adrenocortical transition termination factor 1	2.28
ubiquitin-cytochrome c reductase hinge protein	2.28
scaffold attachment factor 8	2.28
hypothetical protein MG2840 similar to a putative glucosyltransferase	2.28
calumenin	2.27
sialyltransferase 7D ((alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase)	2.27
potassium channel, subfamily K, member 1 (TWIK-1)	2.27
Rho GDP dissociation inhibitor (GDI) alpha	2.27
RNA binding motif protein, X chromosome	2.27
defender against cell death 1	2.27
nuo3 homolog 3 (E. coli)	2.27
polymerase (RNA) III (DNA directed) (G240)	2.27
ubiquitin A-S2 residue ribosomal protein fusion product 1	2.27
tubulin-specific chaperone d	2.27
LIM domain only 2 (Hombobio-Ela 1)	2.26
clathrin, light polypeptide (Lc3)	2.26
protein kinase C beta 1	2.26
putative dimethyladenosine transferase	2.26
chromosome 6 open reading frame 11	2.26
protein phosphatase 6, catalytic subunit	2.26
phosphoribosylglycylamide formyltransferase, phosphoribosylglycylamide synthetase, phosphoribosylaminimidazole synthetase	2.25
hypothetical protein MG2861	2.25
Consensus includes gb:AL096428 /DEF=Human DNA sequence from clone RPS-963E22 on chromosome 20. Contains the 3' end of a novel gene similar to NY-REN-2 Antigen, 5' Cdc islands, E. AL096428	2.25
RNA0071 protein	2.25
dCMP deaminase	2.25
adenylylase kinase 2	2.25
methylenetetrahydrofolate synthetase 2 (beta1)	2.25
hypothetical protein	2.25
M-phase phosphoprotein 9	2.25
RNA0077 protein	2.25
activity-dependent neuroprotector	2.25
serine protease inhibitor, Kunitz type, 2	2.25
clathrin, light polypeptide (Lc4)	2.25
Consensus includes gb:AF062978 /FEA=EST /DB_XREF=gi:5755591 /DB_XREF=est:W2505.x1 /CLONE=HAGE2508513 /UG=Hs.288908 WAS protein family, member 2 /FI=gb:NM_00659 /AP062978	2.24
interleukin 4 receptor	2.24
fragile X mental retardation, autosomal homolog 1	2.24
CD48 antigen (B-cell membrane protein)	2.24
SNR4 structural maintenance of chromosomes 4-like 1 (yeast)	2.24
E1H-domain containing 1	2.24
aryl hydrocarbon receptor interacting protein	2.24
POU domain, class 2, associating factor 1	2.24
ECMT	2.24
Consensus includes gb:AU145293 /FEA=EST /DB_XREF=gi:11006814 /DB_XREF=est:AU145293 /CLONE=HEHMA1004405 /UG=Hs.61773 Homo sapiens cDNA FLJ11648 fs, clone HEHMA1 AU145293	2.24
phosphoglycerate dehydrogenase	2.24
calreticulin	2.23